Tangentially Migrating Transient Glutamatergic Neurons Control Neurogenesis and Maintenance of Cerebral Cortical Progenitor Pools

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The relative contribution of intrinsic and extrinsic cues in the regulation of cortical neurogenesis remains a crucial challenge in developmental neurobiology. We previously reported that a transient population of glutamatergic neurons, the cortical plate (CP) transient neurons, migrates from the ventral pallium (VP) over long distances and participate in neocortical development. Here, we show that the genetic ablation of this population leads to a reduction in the number of cortical neurons especially fated to superficial layers. These defects result from precocious neurogenesis followed by a depletion of the progenitor pools. Notably, these changes progress from caudolateral to rostradorsal pallial territories between E12.5 and E14.5 along the expected trajectory of the ablated cells. Conversely, we describe enhanced proliferation resulting in an increase in the number of cortical neurons in the Gsx2 mutants which present an expansion of the VP and a higher number of CP transient neurons migrating into the pallium. Our findings indicate that these neurons act to maintain the proliferative state of neocortical progenitors and delay differentiation during their migration from extraneocortical regions and, thus, participate in the extrinsic control of cortical neuronal numbers.

**Keywords:** neocortical development, neurogenesis, neuronal numbers, progenitors, tangential migration, transient glutamatergic neurons

Introduction

The generation of an appropriate number of neurons in the central nervous system relies on a strict balance between onset of neurogenesis and maintenance of the progenitor pool (Caviness et al. 2003; Guillemot et al. 2006). At early stages of development, neuroepithelial and radial glia (RG) progenitors are restricted to the ventricular zone (VZ) and divide symmetrically at the apical side to expand the progenitor pool (Gotz and Barde 2005). At the onset of neurogenesis, VZ progenitors begin dividing asymmetrically to generate postmitotic neurons as well as intermediate progenitors (IPs or basal progenitors), which undergo a limited number of divisions in the basal VZ/subventricular zone (SVZ) and, thus, contribute to amplify the number of cortical neurons (Gotz and Huttner 2005; Pontious et al. 2008). Thereafter, newly born neurons migrate radially from the VZ/SVZ to organize in a laminated structure according to an "inside out" sequence whereby early-born neurons reside in deep layers and late-born neurons in more superficial layers (Berry et al. 1964; Bystron et al. 2008). Although the progenitors in the VZ and SVZ undergo both proliferative and neurogenic divisions, their relative contribution to the final number of cortical neurons in deep and superficial layers is still controversial (Pontious et al. 2008; Kowalczyk et al. 2009; Puzzolo and Mallamaci 2010). Nevertheless, the control of their mode of divisions is crucial for the generation of an appropriate number of neuronal subtypes.

An increasing number of molecules regulating cell cycle properties, either the length of specific phases or the timing of cell cycle exit, have been shown to directly influence the mode of divisions of cortical progenitors (Dehay and Kennedy 2007; Glickstein et al. 2009; Puzzolo and Mallamaci 2010). Notably, several of the transcription factors involved in area patterning have also been implicated in the control of cell cycle regulators providing an intrinsic control of progenitors’ proliferation (Muzzio et al. 2005; Holm et al. 2007; Faedo et al. 2008; O’Leary and Sahara 2008). In addition, recent evidence suggests that the secretion of growth factors and morphogens by postmitotic neurons, some of which reach the developing cortex by tangential migration, or by ingrowing thalamic axons provide an extrinsic control for cortical development (Dehay and Kennedy 2007; Komada et al. 2008; Seuntjens et al. 2009; Griveau et al. 2010). However, the mechanisms coordinating intrinsic versus extrinsic processes and their relative contribution to the final cortical cytoarchitecture remain elusive.

We have previously identified a novel population of transient glutamatergic neurons generated by Dbx1-expressing progenitors in the ventral pallium (VP) at the pallial/subpallial boundary (PSB) at E12.5 which progressively invade the pallium by tangential migration to distribute homogeneously in the neonatal cortex (Teissier et al. 2010). We showed that specific genetic ablation of these VP-derived cortical plate (CP) transient neurons, using E1-Ngn2/CRE;Dbx1\(^{DP74}\) animals, leads to a decrease in cortical thickness. In this manuscript, we report that ablation results in a reduced number of cortical neurons which affects predominantly superficial layers and is due to a precocious differentiation and a progressive depletion of the progenitor pools. Conversely, we also describe that Gsx2 mutants, which exhibit an expanded VP (Yun et al. 2001; Stenman, Wang, et al. 2003), display an enhancement in pro-progenitors proliferation as well as in the number of cortical neurons generated correlating with an increase in the number of the CP transient neurons. Our results show that these neurons contribute in a non-cell autonomous manner to maintain the neocortical progenitor pools and, thus, participate in the fine-tuning of cortical neuronal numbers.
Data Collection
Counts of cells labeled by immunofluorescence and measures of the thickness of the deep and superficial layers were performed manually on calibrated pictures using ImageJ software.

1. rD and cl. levels determination: At embryonic stages, the rostro-dorsal (rD) level was identified as the most dorsal domain on coronal sections selected by the presence of the lateral ganglionic eminence (LGE) and septum but not the medial ganglionic eminence. The caudolateral (cl.) level was identified as the domain located just above the morphological hinge at the PSB on coronal sections selected by the presence of the caudal ganglionic eminence and the cortical hem. At E18.5 and P2 stages, the rD level was identified at the level of the prospective primary motor area on coronal sections, whereas the cl. level was identified at the level of the prospective secondary sensory area, according to the Paxinos Mouse Atlases. For each experiment, the sections from the E1-Ngn2/CRE;Dbx1+/TA or Gsx2-/+ mutants were compared with sections from control littermates and processed simultaneously.

2. Measures of deep and superficial layers: The thickness of deep and superficial layers, in the P2 E1-Ngn2/CRE;Dbx1+/TA and control animals, was measured on sections stained by Nissl (Fig. 1A). The layers of the CP were delineated based on the differential cell density. Since the Gsx2-/+ animals do not survive at birth and Nissl staining is not precise enough to delineate cortical layers at E18.5, the thickness of the deep and superficial layers in the E18.5 Gsx2-/+ mutant and control littermates were measured based on coronal sections processed for in situ hybridization for Tbr1 mRNA (Fig. 8A). In all cases, the CP thickness of deep and superficial layers was measured in dorsal and lateral domains of 3 coronal sections for each specimen (only quantifications for rostromedial and caudolateral domains are shown in Figs 1A and 8A).

3. PH3 cells quantifications: For quantifications of VZ and SVZ mitosis at E12.5, E14.5, and E16.5 (Figs 3 and 7), the number of PH3- cells was counted through the thickness of the pallium and normalized for 100-μm-wide strips at both rD and cl. levels on coronal sections. The number of PH3- cells in the VZ corresponds to labeled mitosis located at the border of the ventricle and that for the SVZ corresponds to the abventricular ones.

4. Tbr2 cells quantifications: Tbr2- cells were counted through the thickness of the pallium and normalized for 100-μm-wide strips at both rD and cl. levels on coronal sections (Fig. 5).

5. VZ cell density: The cell density of the VZ was estimated on coronal sections of 14 μm thickness in E16.5 E1-Ngn2/CRE;Dbx1+/TA mutant and control animals by counting the number of DAPI+ cells in 100×100 μm boxes at dorsal and lateral levels. At least 5 boxes were counted for each level in 3 animals.

6. Quantification of the increase in CP transient neurons: The Dbx1Cre, Gsx2Cre;GagCat(CC)-EGFP mice in the Gsx2+/TA and Gsx2-/+ control backgrounds were generated by crossing Gsx2+/TA; GagCat(CC)-EGFP with Gsx2+/TA; Dbx1Cre mice. The total number of GFP+Tbr1- and GFP+Mez2- cells in the CP and MZ (corresponding to CR cells) of the neocortex were counted on coronal sections in 3 specimens for each background.

IdU, BrdU Injections, and Cell Cycle Analysis
Both IdU and BrdU are thymidine analogues which are incorporated during the S-phase of the cell cycle with a clearance in the brain of ~3 h. If we assume that progenitor cells in the telencephalic neuroepithelium 1) progress through the cell cycle asynchronously and 2) have similar cell cycle durations, we can conclude that in such a population of proliferating cells, the fraction of cells in a given phase of the cell cycle is directly proportional to the length of that phase relative to the total length of the cell cycle (Nowakowski et al. 1989). Therefore, upon one single IdU or BrdU injection, all the cells in S-phase at that time will be labeled, and therefore, their proportion among progenitors represents Ts/Tc. If we wait for a ΔT < 3 h for the mice to be sacrificed, the number of labeled cells will therefore represent the proportion of cells that have been in S-phase not only at the time of injections but also during the ΔT and therefore allow to estimate (Ts + ΔT)/Tc (Nowakowski et al. 1989).
Cell Cycle Length

Pregnant mice were injected intraperitoneally with IdU (60 mg/kg), followed by a BrdU injection (50 mg/kg) 1.5 h later and were sacrificed 3 h after the IdU injection to collect the embryos at E12.5 or E14.5. Considering that some DAPI− cells located in the VZ may be negligible as compared with the total number of DAPI+ cells at all levels and stages studied (E12.5: rD = 5.65% in control vs. 5.8% in mutant animals; cL = 7.13% vs. 11.26%; E14.5: rD = 9.42% vs. 9.39%; cL = 9.73 vs. 12.25%; n = 2). In addition, previous studies have shown that a prolonged pulse of BrdU will label virtually all VZ cells at E12.5 (Estivill-Torrus et al. 2002), allowing us to assume that all the DAPI− cells in the VZ were proliferating (Pvcells). Cells that have been labeled upon the IdU injection and that have exited the cell cycle at the time of the BrdU injection will be labeled by IdU and not by BrdU, and their proportion among the total number of proliferative cells is considered as the cells that have exited the cell cycle during one cycle and their proportion among the total number of proliferative cells (IdU− or BrdU+) is designated as the Q fraction (Takehashi et al. 1996; Tarui et al. 2005).

Statistical Analysis

For all experiments, results have been obtained from at least 3 pairs of control and mutant littermates, and the number of counted cells is indicated in the Figure Captions. For all quantifications, normal distribution was confirmed and unpaired, 2-tailed t-test on group indicated in the Figure Captions. For all quantifications, normal distribution was confirmed and unpaired, 2-tailed t-test on group means was performed for statistical analysis, using Microsoft Excel software (* P < 0.05, ** P < 0.005).

Image Acquisition

Brightfield images of brain sections were acquired using a color camera (Zeiss AxioCam HRC) coupled to a Zeiss Axiosvert 200 microscope and immunofluorescence images using an inverted confocal microscope (Leica TCS SP5 AOBS Tandem resonant Scanner).
Results

Reduction of Cortical Neurons’ Numbers Especially in Superficial Layers upon CP Transient Neurons Ablation

We have previously shown that the specific ablation of the CP transient neurons derived from Dbx1-expressing progenitors at the VP/PSB in E1-Ngn2/Cre(iresGFP);Dbx1DTA mutant animals leads to a ~20% reduction in CP thickness throughout the neocortex at E18.5, without affecting either cell density or death (Teissier et al. 2010). This population invades the preplate (PP) and the SVZ/intermediate zone (IZ) by tangential migration with a caudaltonormal to rostromedial gradient starting at E12.5 and is redistributed homogeneously in the CP along the rostrocaudal (RC) and mediolateral (ML) axes at birth.

In order to understand how the loss of Dbx1-derived CP transient neurons leads to a reduction of the CP thickness in E1-Ngn2/Cre(iresGFP);Dbx1DTA animals, we analyzed the defects at caudalateral (cL) and rostrocaudal (rD) levels which represent cortical territories at short and long distances from their generation site, respectively (Teissier et al. 2010). Nissl staining analysis at P2 confirmed that the entire cortex showed a decrease in CP thickness, as previously described at E18.5, with respect to wild-type cortices (rD: 24.32% and cL: 23.87%) (Teissier et al. 2010 and data not shown) and also revealed that this reduction was clearly more pronounced for superficial layers (Fig. 1, rD: 38.98% and cL: 35.77%) than for deep layers (rD: 17.38% and cL: 16.04%) at both rD and cL levels. No significant differences were observed in the extent of the defects between cL and rD levels. These results were confirmed by specifically labeling deep and superficial layers using Tbr1 and Cux1/2 immunostaining, respectively. We observed in both cL and rD regions a small decrease in the thickness of Tbr1+ deep layers in addition to a severe reduction in that of Cux1/2+ superficial layers in mutant brains (Fig. 1B–E). Laminar fate and positioning of cortical neurons appeared to be preserved as observed by immunohistochemistry for FoxP2 (deep layers) and Ctip2 (prospective layer V subpopulations) (Fig. 1F–I) as well as in situ hybridization for Cdh8 (progressive layers II–IV and subpopulation of layer V) and Rorb (progressive layer IV) (Fig. 1J–M). In addition, an increase in the number of FoxP2/labeling was detected in deeper portions of layer VI (Fig. 1F, white bars), whereas that of neurons fated to layer V, as identified by Ctip2 labeling (Fig. 1H), was slightly decreased in mutant compared with control littermates. A more pronounced decrement of Cdh8 and Rorb staining confirmed that superficial layers are more highly affected as revealed by Cux1/2 labeling (Fig. 1J–M). Finally, since another population of tangentially migrating cells, namely Cajal–Retzius (CR) cells, are also generated from Dbx1-expressing progenitors at the VP prior to the CP transient neurons and have been shown to influence cortical patterning (Griveau et al. 2010), we performed whole-mount in situ hybridization for Lmo4 and Cdh8 mRNAs on P0 animals (Supplementary Fig. S1). We observed no differences in area patterning between ablated and control littermates cortices as expected from the absence of defects in CR cells distribution previously reported in the E1-Ngn2/Cre;Dbx1DTA mutants (Teissier et al. 2010) and the late migration and homogeneous distribution of the CP transient neurons.

We conclude that the specific genetic ablation of the CP transient glutamatergic neurons generated from Dbx1-expressing progenitors in the VP leads to a homogeneous deficit in the total number of neurons, although this is more pronounced for the superficial layers, throughout the early postnatal neocortex.

Defects in Differentiation and Proliferation Progressively Affect the Entire Pallium along the Expected Trajectory of the Ablated CP Transient Neurons

Since the CP transient neurons progressively invade the pallium from caudalateral regions starting at E12.0 and reach the rostrocaudal territories by E14.5 (Teissier et al. 2010), we began by analyzing neuronal differentiation at E12.5 and E14.5 using immunostaining for TuJ1 and Tbr1, markers of young postmitotic neurons (Memberg and Hall 1995) and early differentiated glutamatergic cortical neurons (Englund et al. 2005), respectively. We observed an increase in the number of both Tbr1+ and TuJ1+ differentiating neurons in the CP (Fig. 2E–H, black bars) together with ectopically positioned labeled cells in the VZ (arrowheads) in cL and rD regions of E12.5 ablated telencephalons with respect to control littermates. No defects in differentiation were observed in rD regions at this stage (Fig. 2A–D) correlating with the presence of numerous Dbx1-derived CP transient neurons in cL but not in rD territories at E12.5 (Teissier et al. 2010). Interestingly, by E14.5, a similar increase in differentiation was observed in the rD region of mutant telencephalons (Fig. 2F and data not shown), whereas no differences were detected in the CP of the cL region, with the exception of ectopic Tbr1+ neurons in the VZ (Fig. 2MN, arrowheads). Moreover, at E14.5 a slight thinning of the VZ was apparent especially in mutant cL territories (Fig. 2MN, black bars).

In order to assess if the increase in differentiation was due to defects in proliferation occurring upon ablation, we quantified the number of mitotic cells, as labeled by PH3 staining, in the rD and cL regions of the developing pallium at E12.5, E14.5, and E16.5 (Fig. 3A–D). We detected an initial increase in the number of PH3+ mitotic progenitors in the cL VZ at E12.5 followed by a reduction clearly observed at E16.5 (Fig. 3C). At E14.5, corresponding to the time of arrival of the CP transient neurons in rD territories in control animals, enhanced VZ proliferation was observed in rD regions of mutant cortices (Fig. 3AEF), followed by a small decrease by E16.5 (Fig. 3A). Notably, the number of PH3+ mitotic progenitors in the SVZ was strongly reduced at both cL and rD levels starting at E14.5 (Fig. 3BDEH).

Taken together, these results show that the loss of the CP transient neurons leads initially to an increment in the number of mitosis in the VZ and of differentiating neurons (at E12.5 in cL and E14.5 in rD regions), followed by a decrement in the number of both VZ and SVZ mitosis. Notably, changes in proliferation and differentiation appear to progress from the caudalateral to the rostrocaudal cortex and to correlate with the timing and trajectory of invasion of pallial regions by the CP transient neurons.

CP Transient Neurons Ablation Results in Precocious Neurogenesis

The defects observed in ablated animals could result from either a change in cell cycle length, an increase in the fraction of neurogenic divisions and/or a variation in the size of the progenitor pools. In order to discriminate between these possibilities, we started by measuring the length of the cell cycle using IdU and BrdU injections 3 and 1.5 h before collection of the embryos (see Materials and Methods) (Fig. 4A–C). At E12.5, we observed lengthening of the cell cycle in progenitors located in the VZ in cL (Tc = 13.31 h ±
2.91 in controls and Tc = 25.71 h ± 4.02 in mutants) but not in rD territories (Tc = 9.19 h ± 1.07 in controls and Tc = 11.96 h ± 1.09 in mutants). By E14.5, VZ progenitors at both cl and rD levels displayed a clear increase in Tc (rD: Tc = 11.74 h ± 0.54 in controls and Tc = 18.51 h ± 3.27 in mutants; cl: Tc = 17.81 h ± 2.80 in controls and Tc = 27.96 h ± 3.91 in mutants). Therefore, defects in proliferation and differentiation correlate with a lengthening of the cell cycle in E1-Ngn2/Cre;Dbx1DTA mutant embryos.

A lengthening of the cell cycle has been shown to characterize differentiative divisions (Miyama et al. 1997; Calegari et al. 2005). To address whether an increased fraction of cells exited the cell cycle (Q fraction) in E14.5 E1-Ngn2/Cre;Dbx1DTA animals, we performed a single injection of IdU followed by multiple injections of

Figure 2. Transient increase in differentiation progressively affects the entire pallium of E1-Ngn2/Cre;Dbx1DTA mutants. (A–H) Immunohistochemistry performed on E12.5 embryos shows no differences for Tbr1 (A,B) and TuJ1 (C,D) staining between control (A,C) and E1-Ngn2/Cre;Dbx1DTA (B,D) embryos at rD levels, whereas there is a strong increase for both in cl regions in mutants (F,H, black bars) compared with control littersmates (E,G). Moreover, ectopic Tbr1⁺ and TuJ1⁺ cells are observed in the mutant VZ (arrowheads in F,H). (I–N) By E14.5, Tbr1 staining is increased at rD levels in mutants (J,L), whereas no differences are detected at cl levels (I,K) with respect to control littersmates (I,K,M). Ectopic Tbr1⁺ neurons (arrowheads in I,L) are observed in the VZ of mutants at both levels as well as a thinning of the VZ at cl levels (black bars in M,N). K and L represent high magnifications of boxed areas in I and J, respectively. Scale bars: A,K, 20 μm; I, 100 μm.
levels, both VZ and SVZ display reduced number of mitosis in mutant animals (Immunostaining for PH3 at both rD (Fig. 4A,B) and cL (CD) levels at E12.5, E14.5, and E16.5. Quantifications of the number of PH3+ cells show a transient increase in VZ proliferation of cl regions at E12.5 (P = 0.02402; n = 3) and of rD regions at E14.5 (P = 0.02487; n = 4) followed by a decrease at all levels by E16.5 (rD: P = 0.15084; cl: P = 0.0017; n = 4). A reduction of SVZ proliferation is detected at E14.5 and E16.5 at both levels (rD: E14.5, P = 0.01482 and E16.5, P = 0.03273; cl: E14.5, P = 0.03167 and E16.5, P = 0.00100). Graphs represent means ± SEM. *P < 0.05 and **P < 0.005. (E-H) Immunostaining for PH3 at both rD (E,F) and cl (G,H) levels at E14.5 shows more numerous mitoses in the VZ with fewer mitoses in the SVZ of mutants (F) at rD levels. At cl levels, both VZ and SVZ display reduced number of mitosis in mutant animals (H) compared with controls (G). Scale bars: E, 20 μm.

Brdu (every 2 h) for a total of 15 h (Fig. 4D) and quantified the number of cells which failed to reentered the S-phase during the time of the experiment (IdU+Brdu+) among actively cycling cells (IdU+Brdu+ or IdU+Brdu–) (see Materials and Methods). We observed that an increased proportion of cells exited the cell cycle in ablated cortices compared with controls at both cl and rD levels (Fig. 4E). Evaluation of the total number of postmitotic neurons generated during one cell cycle by counting the number of Tbr1+ cells relative to the size of the progenitor pool (IdU+BrdU+) also confirmed an enhanced fraction of neurogenic divisions in mutant compared with control littersmates (Fig. 4F).

Furthermore, we detected an increase in the expression of Ngn2, a proneural gene detected at high levels in neurogenic divisions (Shimojo et al. 2008; Ochiai et al. 2009), in the VZ and SVZ of mutant compared with control littersmates (Fig. 4G,H). Together, these experiments suggested that there was an augmentation in the number of progenitors undergoing neurogenic instead of proliferative divisions in both the VZ and SVZ of ablated embryos.

To further analyze whether the cortical neuroepithelium was undergoing precocious neurogenesis, we performed immunostaining for Tbr2, a gene expressed in both IP cells and early postmitotic neurons in the developing pallium (Englund et al. 2005) and TuJ1. We observed an increase in the number of TuJ1+ cells in the VZ of E14.5 E1-Ngn2/Cre(iresGFP);Dbx1DTA animals throughout the neocortical neuroepithelium (Englund et al. 2005; TuJ1+ cells in the apical VZ, but not reciprocally, in both control and mutant E1-Ngn2/Cre(iresGFP);Dbx1DTA cortices (Fig. 4K-P), suggesting that the GFP labels a specific subpopulation of the Tbr2+ cells. Moreover, the TuJ1+Tbr2+ cells in the apical VZ observed in control as well as the ectopic ones observed in mutant animals were almost always associated with GFP expression (green arrowheads in Fig. 4K–R), suggesting that the Tbr2+GFP+ cells in the apical VZ are postmitotic neurons. Accordingly, we observed a higher proportion of Tbr2+GFP+ cells among Tbr2+ cells at E12.5 in cl territories (cl: 20.6 ± 2.65% in controls compared with 51.4 ± 2.96% in mutants, the GFP is not expressed at rD level at this stage). A similar increase was detected in both cl and rD regions at E14.5 (cl: 31.85 ± 1.79% in controls compared with 52.76 ± 0.18% in mutants; rD: 29.39 ± 11.79% in controls compared with 63.19 ± 3.03% in mutants) and maintained at E16.5 (cl: 15.06 ± 1.40% in controls compared with 45.82 ± 9.68% in mutants; rD: 15.75 ± 0.78% in controls compared with 24.67 ± 0.46% in mutants). We conclude that an enhanced fraction of Tbr2+ cells corresponds to postmitotic neurons upon ablation of the CP transient neurons. Thus, together these experiments demonstrate that progenitors throughout the neocortical neuroepithelium undergo precocious neurogenic divisions in E1-Ngn2/CRE;Dbx1DTA animals by midneurogenesis.
Depletion of Neocortical Progenitor Pools in Ablated Mutants

An increase in the neurogenic fraction of progenitor divisions and a decrease in that of proliferative cells should lead to a progressive depletion of the progenitor pool. Indeed, the observed decrease in the number of mitosis in E14.5 and E16.5 in E1-Ngn2/Cre;Dbx1DTA embryos together with a reduction in Ki67 staining, a marker of proliferative cells, at caudal levels of E14.5 mutant pallium compared with control littermates (Fig. 5A–B) strongly suggested a global depletion of the progenitor pools.

Since the decrease in the proliferation was most pronounced in the SVZ at E14.5, we first tested whether the pool of IPs was affected. We quantified the number of Tbr2+ cells within one column spanning the thickness of the developing pallium and observed a reduction starting in the cl. region of mutant telencephalons at E12.5 (Fig. 5C–E). Strong defects were observed in rD territories at E14.5 and E16.5 but not at E12.5. Therefore, together with the enhanced number of postmitotic Tbr1+ and Tbr2+ neurons at E12.5 and E14.5 in cl. and rD regions, respectively (Figs 2E,F,I–L and 4K–P) and the reduction in SVZ...
proliferation (Fig. 3B,D), these results show that the decrease in the total number of Tbr2\(^+\) cells corresponds to a depletion in the pool of IPs rather than of early postmitotic neurons (also Tbr2\(^+\)).

In order to determine if the pool of RGs was also affected, we analyzed the expression of BLBP and Pax6 proteins, 2 specific markers of RG cells, at E16.5. We observed a decrease in their expression in mutant cL cortices (Fig. 5F,G) correlating with a ~25% reduction in cell density, as measured by DAPI staining, in the VZ (Fig. 5L). In addition, we observed at E16.5 a decrease in BrdU labeling (upon a 2 h pulse) in the VZ and SVZ of both rD and cl regions (Fig. 5J,K). We conclude that a depletion of both the IP and the RG progenitors’ pools occurs in the pallium of the E1-Ngn2/Cre;Dbx1\(^{DTA}\) animals.

Increased Number of Dbx1-Derived CP Transient Neurons in the Pallium of Gsx2 Mutants

Since the E1-Ngn2/CRE;Dbx1\(^{DTA}\) mutants represent a specific ablation of Dbx1-expressing progenitors at the VP, we sought to analyze Gsx2 null mice which have been reported to display opposite effects, namely an expansion of the Dbx1 expression domain in the VP (Yun et al. 2001; Carney et al. 2009). Notably, these mutants have recently been shown to present an increase

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**Figure 5.** Depletion of RG and IP progenitor pools in E1-Ngn2/CRE;Dbx1\(^{DTA}\) animals. (A-B’) Immunohistochemistry for Ki67 shows a global decrease in cycling progenitors in mutant E1-Ngn2/CRE;Dbx1\(^{DTA}\) (B’) compared with control (A,A’) cortices at E14.5. (C,D) Immunohistochemistry for Tbr2 reveals fewer Tbr2\(^+\) cells in the pallium of E14.5 mutant (D) compared with control (C) littersmates. (E) Graphs represent the number of Tbr2\(^+\) cells through the thickness of the pallium and normalized for 100-\(\mu\)m-wide strips at E12.5 at both cl and rD levels and E14.5 and E16.5 at rD levels in controls (black columns) and mutants (white columns). The number of Tbr2\(^+\) cells is reduced, although not statistically significant, at cl levels of E12.5 mutant embryos. At rD levels, a 22% decrease is observed at E14.5 and E16.5 (E14.5: \(P = 0.023539\); E16.5: \(P = 0.023532\)). (F-K) Immunohistochemistry for BLBP (F,G) and Pax6 (H,I) on E16.5 control (F,H) and mutant (G,I) animals shows a reduction in the size of the RG pool in the VZ of mutant animals. (J,K) A pulse of BrdU 2 h before sacrifice labeling cells in S-phase reveals fewer cycling progenitors in both VZ and SVZ of E16.5 mutant (K) compared with control (J) littersmates. (L) Graph shows the number of DAPI\(^+\) nuclei per 100 \(\mu\)m\(^2\) boxes of 14-\(\mu\)m-thick coronal sections in the VZ at dorsal and lateral levels in E16.5 mutants (white columns, dorsal: 211.66 ± 22.98, lateral: 225.87 ± 12.87) and controls (black columns, dorsal: 311.62 ± 8.3, lateral: 313.56 ± 11.43) (\(P = 0.026\) in dorsal and \(P = 0.036\) in lateral). Graphs represent means ± SEM. *\(P < 0.05\). Scale bars: A,C, 100 \(\mu\)m; A’,F,H, 50 \(\mu\)m; and J, 20 \(\mu\)m.
in the number of Dbx1-derived glutamatergic neurons migrating into the lateral amygdala (Waclaw et al. 2010). The role of Gsx2 in dorsoventral patterning and neurogenesis of the subpallium has been analyzed in several studies (Corbin et al. 2000; Yun et al. 2001; Stenman, Toresson, et al. 2003; Yun et al. 2003; Carney et al. 2009). However, no major defects in pallial development had been reported so far.

In order to permanently trace Dbx1-derived CP transient neurons in the pallium of the Gsx2 mutants, we crossed Dbx1\textsuperscript{CRE}\textsuperscript{+};Gsx2\textsuperscript{R26R LacZ} with CagCat-EGFP;Gsx2\textsuperscript{R26R LacZ} animals (Waclaw et al. 2010). By immunostaining for Tbr1 on E18.5; Carney et al. 2009). However, no major defects in pallial

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Enhanced Proliferation and Neurogenesis in the Pallium of Gsx2 Mutants

We started by analyzing the progression of differentiation using Tbr1 and Tuj1 immunostaining in E12.5 Gsx2 mutants and observed a global decline affecting both cl and rD regions, but this was more pronounced at cl levels (Supplementary Fig. S2). We also detected a concomitant enhanced proliferation in both the VZ and the SVZ, as revealed by Ki67 and BrdU staining (Supplementary Fig. S2). At E14.5, a strong increment in Ki67 and Ph3 immunostaining was detected in both the VZ and the SVZ of Gsx2 mutants at cl and rD levels (Fig. 7A–F and data not shown). Quantification of Ph3\textsuperscript{+} cells at the apical VZ and the SVZ confirmed a higher number of ventricular and subventricular mitosis in these mutants (Fig. 7M,N). In addition, we observed more numerous Tbr2\textsuperscript{+} cells in the SVZ (Fig. 7G,H) but not at the apical VZ (Fig. 7G,H,O), showing that both RG and IP progenitors display enhanced proliferation. Finally, we observed a reduction in Ngn2 staining in both the VZ and the SVZ of mutant animals (Fig. 7J) as well as a reproducible slight decrease in Tis21 expression, especially observed in the apical portion of the VZ (Fig. 7K,L), strongly arguing for a decrease in the fraction of neurogenic divisions. Notably, the number of Tbr1\textsuperscript{+} neurons in the CP of E14.5 Gsx2 mutants was similar to that of control embryos (data not shown). Therefore, together these results suggest that an amplification of both RGs and IPs progenitor pools and a delayed neurogenesis occur in the pallium of Gsx2 mutants. These defects are opposite with respect to the precocious differentiation and depletion of the progenitor pools described in the E1-Ngn2\textsuperscript{CRE};Dbx1\textsuperscript{I\textasciitildeTA} mutants at corresponding stages.

We then analyzed the phenotype at E18.5, the latest stage possible for Gsx2 mutants which die at birth and observed an increase in the thickness of the CP at both rD (25.49%) and cl (30.96%) levels. Tbr1 and Cux1\textasciitilde2 immunostaining showed that both deep and superficial layers were thicker (Fig. 8B–G) and appeared to be affected in a similar manner (Fig. 8A), although the supernumerary CP transient neurons were positioned in deep layers. Finally, we observed a strong increase of neuronal numbers in layers V and II–IV upon immunostaining for FoxP2, Ctip2 and Mef2C, as well as by in situ hybridization for Calb2 (Fig. 8H–I). In conclusion, we have shown that Gsx2 mutants present a surplus number of CP transient neurons and display opposite non-cell

![Figure 6](https://example.com/figure6.png)
autonomous pallial defects to those observed upon their ablation in E1-Ngn2/CRE;Dbx1<sup>DTA</sup> animals, namely excessive proliferation of both VZ and SVZ progenitors and a consequent increased number of cortical neurons fated to both deep and superficial layers.

**Discussion**

We have shown that the specific ablation of CP transient neurons derived from Dbx1-expressing progenitors results in a decrement in the final number of cortical neurons as a consequence of the precocious differentiation of cortical progenitors. The kinetics of the defects observed in ablated mutants follow the expected trajectory of these cells, which migrate tangentially from the VP at the border of the developing pallium (Teissier et al. 2010). On the contrary, a gain of CP transient neurons, as observed in Gsx2 mutants, correlates with a delayed onset of neurogenesis in the pallium resulting in an increase in the size of the progenitor pools and in the total number of cortical neurons. Therefore, all together these results show that these transient glutamatergic neurons from the postmitotic compartment appear to control the mode of divisions of cortical progenitors in a non-cell autonomous manner over long distances from their generation site. Their dynamics of invasion of the CP together with their role in delaying differentiation might represent a novel mechanism to refine neurogenetic gradients during cortical growth via cell migration.

**The Proliferation/Differentiation Transition of Cortical Progenitors Depends on the Number of CP Transient Neurons in Developing Pallial Territories**

The specific ablation of Dbx1-derived CP transient neurons results in an early increase in the leaving fraction and a lengthening of the cell cycle together with an enhanced expression of markers of neurogenic divisions. Correlating with the observed precocious neurogenesis, the generation of FoxP2<sup>+</sup> layer VI neurons is incremented in E1-Ngn2/CRE;Dbx1<sup>DTA</sup> animals. In addition, proliferation in the SVZ is reduced suggesting a defect in IPS generation from VZ progenitors. Thus, the early onset of differentiation leads to a progressive depletion of both VZ and SVZ progenitor pools and results in a general reduction of neuronal production although this is more pronounced for neurons fated to superficial layers. On the contrary, in Gsx2 mutants, which display an excessive generation of VP-derived CP transient neurons, the numbers of cycling VZ and SVZ progenitors and of cortical neurons are amplified. Thus, a delayed differentiation resulting in an enlarged progenitor pool occurs in Gsx2 mutants. However, unlike E1-Ngn2/CRE;Dbx1<sup>DTA</sup> mutants, Gsx2 mutants display a fairly homogeneous increase of both superficial and deep layer neurons. This is likely due to the fact that E1-Ngn2/CRE;Dbx1<sup>DTA</sup> mutants present only a partial (around 50%) ablation of the CP transient neurons (Teissier et al. 2010) because of a late onset of recombination starting after E11.5 whereas VP expansion is already well established at this stage in Gsx2 mutants (Yun et al. 2001). Nevertheless, our results show that E1-Ngn2/CRE;Dbx1<sup>DTA</sup> and Gsx2 mutants represent mouse models for studying loss and gain of function of Dbx1-derived CP transient neurons, respectively. In addition, this is the first time that a pallial phenotype is described for Gsx2 mutants. Indeed, previous studies have mostly focused on the
role of Gsx2 on dorsoventral patterning and have described a considerably smaller stratum resulting from patterning defects that lead to a specific decrease in proliferation in the LGE, notably affecting the SVZ at E12.5 (Corbin et al. 2000; Toresson et al. 2000; Toresson and Campbell 2001). This cell autonomous phenotype appears opposite to the one we describe in the pallium and supports an indirect function for Gsx2 in the development of the cortical primordium through the inhibition of expression of the Dbx1 gene (Yun et al. 2001; Winterbottom et al. 2010).

All together, our results strongly suggest that CP transient neurons have an active role in enhancing progenitor proliferative divisions and, thus, decreasing the probability of cell cycle exit. In addition, both E1-Ngn2/CRE/Dbx1Dota and Gsx2 mutants are strongly affected in SVZ proliferation and, present defects in both deep and superficial layers, supporting the hypothesis that IPs generate neurons fated to all layers and have a role in the amplification of RGs progeny rather than on neuronal fate (Pontious et al. 2008; Kowalczyk et al. 2009; Puzzolo and Mallamaci 2010). Accordingly, both CyclinD2 and Tbr2 mutants specifically affecting SVZ proliferation display a reduced number of neurons in both deep and superficial layers (Arnold et al. 2008; Sessa et al. 2008; Glickstein et al. 2009) although this is more pronounced in superficial layers. However, this appears to be in the absence of precocious differentiation of VZ progenitors. On the contrary, mutants that do result in premature differentiation, such as Pax6 and Tlx mutants, also display a depletion of the progenitor pool correlated with an increased leaving fraction and a consequent reduction in the thickness of superficial layers as described for E1-Ngn2/CRE/Dbx1Dota animals (Land and Monaghan 2003; Roy et al. 2004; Tuoc et al. 2009; Georgala et al. 2011). Interestingly, mutants for both Pax6 (Sey/Sey) and Tlx present a loss of Dbx1 expression at the VP/PSB (Yun et al. 2001; Stenman, Yu, et al. 2003; Roy et al. 2004), and cell death has been reported to specifically occur at the PSB of Tlx mutants (Roy et al. 2004). Although both the Pax6 and the Tlx genes have been shown to control cell autonomously the mode of divisions of cortical progenitors (Quinn et al. 2007; Osumi et al. 2008; Zhang et al. 2008) it cannot be excluded that part of the phenotype in their respective mutants is dependent on alterations in the generation or specification of the CP transient neurons due to VP defects.

**CP Transient Neurons Migration and “Neurogenic Gradients”**

Gradients of proliferation (rostrolateral\textsuperscript{high} to caudomedial\textsuperscript{low}) and differentiation (lateral\textsuperscript{high} to medial\textsuperscript{low}) have been suggested to regulate timing of early neurogenesis and, thus, neuronal numbers in distinct cortical territories (Bayer et al. 2001; 2003; 2004). Notably, both deep and superficial layers appear to be thickened in the mutant cortices in a similar manner (rD: P < 0.001; cL: P = 0.0121). Notably, both deep and superficial layers appear to be thickened in the mutant cortices in a similar manner (rD: P < 0.001; cL: P = 0.0121). We controlled that the thickening of cortical layers was not the consequence of a reduction in neuronal density by counting the number of DAPI\textsuperscript{+} nuclei cells in the CP and we rather observed the opposite effects (120 ± 4.82\% mutant/control, n = 3), therefore suggesting that the defects in thickness were likely underestimated. Graphs represents mean ± SEM. *P < 0.05. (A–G) Immunostaining for Tbr1 (B–F) and Cux1/2 (F,G) labeling deep and superficial layers, respectively, confirms a thicker CP comprising both the deep and the superficial layers in E18.5 Gsx2\textsuperscript{mutant} (C,E,G) compared with Gsx2\textsuperscript{+/+} control (B,D,F) littersmates. (H–I) Immunostaining at rD levels for FoxP2 (H,J), Ctip2 (J,K), and Mef2c (K,L) and in situ hybridization for Cdh8 (L,M), labeling layers V/VI, layer V, and layers II–IV. Except for neurons located in layer VI whose number appears reduced (white bars in H,J), all other markers show a pronounced increase in Gsx2 mutants (I,K,O) compared with controls (H,J,L,N). Scale bars: B, 200 μm; D,H,L, 100 μm.

**Figure 8.** E18.5 Gsx2 mutants present an increase in the thickness of both deep and superficial cortical layers. (A) Graphs represent the thickness of deep (white) and superficial (gray) layers in Gsx2 mutants relative to the thickness of Gsx2\textsuperscript{+/+} control animals in rostrodorsal and caudolateral matched territories based on Tbr1 mRNA differential expression in deep versus superficial layers (rD: deep layers: 128.68\%, P = 0.0478; superficial layers: 122.45\%, P = 0.0122; n = 5 and cL: deep layers: 150.13\%, P = 0.0146; superficial layers: 120.19\%, P = 0.0121). Notably, both deep and superficial layers appear to be thickened in the mutant cortices in a similar manner (rD: P < 0.001; cL: P = 0.0120). We controlled that the thickening of cortical layers was not the consequence of a reduction in neuronal density by counting the number of DAPI\textsuperscript{+} nuclei cells in the CP and we rather observed the opposite effects (120 ± 4.82\% mutant/control, n = 3), therefore suggesting that the defects in thickness were likely underestimated. Graphs represents mean ± SEM. *P < 0.05. (A–G) Immunostaining for Tbr1 (B–F) and Cux1/2 (F,G) labeling deep and superficial layers, respectively, confirms a thicker CP comprising both the deep and the superficial layers in E18.5 Gsx2\textsuperscript{mutant} (C,E,G) compared with Gsx2\textsuperscript{+/+} control (B,D,F) littersmates. (H–I) Immunostaining at rD levels for FoxP2 (H,J), Ctip2 (J,K), and Mef2c (K,L) and in situ hybridization for Cdh8 (L,M), labeling layers V/VI, layer V, and layers II–IV. Except for neurons located in layer VI whose number appears reduced (white bars in H,J), all other markers show a pronounced increase in Gsx2 mutants (I,K,O) compared with controls (H,J,L,N). Scale bars: B, 200 μm; D,H,L, 100 μm.
maintenance of RGs proliferation through the activation of Notch (Caviness et al. 2009; Liu et al. 2010). In addition, newly generated molecules secreted by the patterning centers and to progress by cell–cell contact through hemichannels propagating calcium waves. This process has been shown to be both necessary and sufficient to promote RGs proliferation (Bittman et al. 1997; Takahashi et al. 1999). We have shown that the CP transient neurons are generated with a peak at E12.5 and migrate tangentially to reach the rostrodorsal pallium by E14.5 along caudal to rostral and lateral to dorsal migratory pathways (Teissier et al. 2010). Therefore, CP transient neurons’ migration parallels the neurogenic gradient occurring between E12.5 and E14.5. Strikingly, while the defects in proliferation follow the expected trajectory of ablated cells in E1-Ngn2/CRE;Dbx1IPTA animals (i.e., starting at the caudal PSB at E12.0–E12.5 and reaching the dorsal cortex by E14.5), the decrease in neuronal generation is homogeneous along the RC axis despite delayed onset of ablation effects in rostrodorsal regions, suggesting that the neurogenic gradients are unperturbed in ablated animals. Since the pattern of the E1/Ngn2 enhancer activity also correlates with gradients of differentiation (Berger et al. 2004) and the expression of Ngn2 increases in mutant animals, our data strongly suggest that ablation results in a switch to neurogenic divisions at the expense of proliferative divisions (lacopetti et al. 1999; Kawaguchi et al. 2004). Dbx1-derived CP transient neurons might act by delaying differentiation possibly by maintaining low Ngn2 expression in progenitors. This occurs in the absence of defects in laminar fate and arealization, strongly suggesting that CP transient neurons do not affect the intrinsic sequence which leads progenitors to produce the whole set of layer-specific neurons once they have initiated neurogenesis (Shen et al. 2006; Gaspard et al. 2008). Most likely, they appear to time the exit from the cell cycle and, thus, to increase the total number of self-renewing cell cycle per progenitor.

**CP Transient Neurons Migration Provides an Extrinsic Mechanism to Control the Mode of Divisions of Cortical Progenitors**

We have shown that both RGs and IPs progenitors are affected upon modifications in the number of CP transient neurons. Our results indicate that the CP transient neurons might influence progenitors proliferation/differentiation transition while invading the pallium. Since these neurons migrate through both the IZ/SVZ and the developing PP at early stages of development (E12.5–E14.5) (Teissier et al. 2010), they could interact at this location by cell-cell-mediated contacts with the basal processes of RG cells or with IPs as well as by the secretion of signaling factors. Notably, the neurogenic gradient observed in the developing cortex has already been suggested to be initiated by signaling molecules secreted by the patterning centers and to progress by cell–cell contact through hemichannels propagating calcium waves. This process has been shown to be both necessary and sufficient to promote RGs proliferation (Bittman et al. 1997; Caviness et al. 2009; Liu et al. 2010). In addition, newly generated neurons and IPs are involved by presenting Delta ligands in maintenance of RGs proliferation through the activation of Notch signaling (Yoon et al. 2008). Lastly, an increasing number of evidence suggests that the secretion of diffusible molecules provides a feedback control from the postmitotic compartment to regulate the proliferation properties of cortical progenitors (Polleux et al. 1998; Dehay et al. 2001; Viti et al. 2003; Agasse et al. 2006; Dehay and Kennedy 2007; Scuettens et al. 2009; Griveau et al. 2010). However, it is unlikely that Dbx1-derived CP transient neurons would use the same molecules expressed by all differentiating glutamergic neurons to communicate with cortical progenitors, for example, Nt3/Fgf9 as in the case of Sip1 mutants (Scuettens et al. 2009) or Notch ligands (Yoon et al. 2008). Indeed, they represent a small percentage of the neurons in the postmitotic compartment at any given stage and especially in rD at E14.5 when the effect upon ablation is already observed (Teissier et al. 2010) and their ablation would further be compensated by the precocious differentiation, initially generating more neurons.

Patterning, growth, and cell fate, by “mobile signaling cells/structures,” such as TCA axons and migrating cells (CR neurons, CP transient neurons, neural crests, meninges, and blood vessels) (Etchevers et al. 1999; Dehay et al. 2001; Schneider et al. 2001; Creuzet et al. 2004; Vasudevan et al. 2008; Javaherian and Kriegstein 2009; Siegenthaler et al. 2009; Stubbbs et al. 2009; Griveau et al. 2010) depend critically on the timing of their generation, migration, and arrival in different cortical regions. Their early arrival (E10.5–E12.5) will affect regionalization and later arrival (E12.5–E13.5) the onset of neurogenesis in a tight interplay with intrinsic developmental programs of cortical progenitors. Therefore, it appears that multiple temporal, spatial, and qualitative levels of control underlie the role of these mobile “signaling” cells/structures in cortical development. The identification of this novel population and its crucial non-cell autonomous role in progenitor pool maintenance will now allow further studies aiming at the molecular characterization of how migrating CP transient neurons contributes to the extrinsic control of corticogenesis.

**Supplementary Material**

Supplementary material can be found at: http://www.cercor.oxfordjournals.org/

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