

Cdk6-Dependent Regulation of G₁ Length Controls Adult Neurogenesis

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ABSTRACT

The presence of neurogenic precursors in the adult mammalian brain is now widely accepted, but the mechanisms coupling their proliferation with the onset of neuronal differentiation remain unknown. Here, we unravel the major contribution of the G₁ regulator cyclin-dependent kinase 6 (Cdk6) to adult neurogenesis. We found that Cdk6 was essential for cell proliferation within the dentate gyrus of the hippocampus and the subventricular zone of the lat-

eral ventricles. Specifically, Cdk6 deficiency prevents the expansion of neuronally committed precursors by lengthening G₁ phase duration, reducing concomitantly the production of newborn neurons. Altogether, our data support G₁ length as an essential regulator of the switch between proliferation and neuronal differentiation in the adult brain and Cdk6 as one intrinsic key molecular regulator of this process. *STEM CELLS* 2011;29:713–724

Disclosure of potential conflicts of interest is found at the end of this article.

INTRODUCTION

Brain plasticity is expressed in diverse manners in adulthood and includes de novo neuronal production. This evolutionary conserved process occurs at least in two specific compartments of the adult brain: the subgranular zone (SGZ) of the hippocampal dentate gyrus (DG) and the subventricular zone (SVZ) of the lateral ventricles whose neural precursor cells (NPC) give rise to dentate granule neurons and olfactory bulb (OB) interneurons, respectively [1]. Although much has been elucidated about the identity and fate of adult precursor cells, the mechanisms that couple proliferation with neuronal differentiation in the adult brain remain largely unknown.

During the cell cycle, G₁ is an essential phase in which there is integration and response to extracellular cues that either allow progression through the cell cycle or promote withdrawal from the cell cycle to embark on a differentiation pathway [2]. Progression through G₁ is driven by the concerted action of cyclin-dependent kinase (Cdk) 4 (Cdk4) and the closely-related Cdk6 and their activating partners, that is, the D-type cyclins [3]. Once activated by D-type cyclins, Cdk4/6 phosphorylate the retinoblastoma protein (pRb), leading pRb to release E2fs. E2fs are transcriptional activators

that, once freed from pRb, are able to activate the transcription of genes necessary to enter S-phase [4, 5]. Conversely, Cdk4/6 activity is suppressed through interactions with members of two families of inhibitory proteins: the Ink4 proteins (p15^{Ink4a}, p16^{Ink4b}, p18^{Ink4c}, and p19^{Ink4d}) that exhibit selectivity for Cdk4/6, and the Cip/Kip proteins (p21^{Cip1}, p27^{Kip1}, and p57^{Kip2}) that have a broader range of Cdk inhibitory activity [6].

Recently, accumulating evidence ascribed crucial roles to regulators of Cdk4/6 activity in controlling adult neurogenesis. For instance, p27^{Kip1} selectively constrains transit-amplifying cell proliferation in the SVZ [7], whereas deletion of Cyclin D2 virtually abrogates adult neurogenesis [8]. However, the direct and specific contribution of the catalytic partners Cdk4 and Cdk6 to adult neurogenesis has not been evaluated yet. Moreover, to date, no study reported how cell cycle and particularly G₁ parameters are affected in adult neural precursors following genetic ablation of G₁ regulators.

In this report, we used knockout mice for the G₁ regulators Cdk4 and Cdk6 to analyze their respective contribution to adult neurogenesis. We showed that Cdk6 specifically controls the expansion of neuronally committed precursors and their rate of cell cycle withdrawal by regulating the length of G₁. Thus, our study describes for the first time the importance

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of regulating G_1 duration for fine-tuning neuronal production in the adult brain.

MATERIALS AND METHODS

Animals

Cdk4^{-/-} [9] and Cdk6^{-/-} [10] colonies were maintained on a mixed CD1 × C57BL/6J × FVB × S129/sv and a mixed CD1 × C57BL/6J background, respectively. Wild type (WT) and knock-out mice used in all experiments were obtained from heterozygous breedings and genotype was determined by polymerase chain reaction as described previously. Unless otherwise mentioned, 6-week-old male and female mice were used in all experiments. Mice were group-housed in the animal facility at the University of Liège under standard conditions with food and water ad libitum and were maintained on a 12-hour light/dark cycle. All animals were taken care of in accordance with the declaration of Helsinki and following the guidelines of the Belgian Ministry of Agriculture in agreement with European Community laboratory Animal Care and Use Regulations (86/609/CEE, Journal Officiel des Communautés Européennes number L358, 18 December 1986).

Seizure Induction

See Supporting Information.

BrdU Administration and Detection

See Supporting Information.

Immunofluorescence

Tissue was prepared as described in Supporting Information. All staining was performed on 40- μ m free-floating sections. Sections underwent antigen retrieval for 30 minutes at 95°C in Target Retrieval Solution (Dako, Glostrup, Denmark), were then rinsed in Tris-buffered saline (TBS) and incubated overnight at 4°C with primary antibodies (complete list of primary antibodies as Supporting Information) diluted in TBS containing 0.1% Triton, 0.1% Tween 20 and 5% normal donkey serum (blocking solution). After washings in TBS, sections were incubated for 1 hour at room temperature in blocking solution containing the corresponding secondary antibodies (1:500, Jackson ImmunoResearch, Europe Ltd, Suffolk, U.K.), either coupled to Fluorescein Isothiocyanate (FITC), Rhodamine Red-X, or Cy5. Finally, sections were rinsed in TBS and mounted between slide and coverslide using VectaShield Hard Set mounting medium containing 4',6-diamidino-2-phenylindole (DAPI) (Vector Laboratories, Burlingame, CA). The slides were stored in the dark at 4°C.

Confocal Microscopy

Fluorescence images were acquired using the Olympus Fluoview FV1000 confocal system equipped with the Olympus IX81 inverted microscope (Olympus Europa GmbH, Hamburg, Germany). Fields were acquired using Z-scan with a step of 1.5 μ m between each confocal plane. All sections prepared for comparison were analyzed at the same time, using the same acquisition parameters.

Quantification of Cell Numbers

All quantifications were realized by an experimenter blind to the experimental conditions. To evaluate cell proliferation and neurogenesis in the DG, we counted exhaustively in the SGZ (SGZ + granule cell layer [GCL] for bromodeoxyuridine [BrdU] 4 weeks analysis), the number of Ki-67⁺ or BrdU⁺ cells per DG section under a 60 \times objective in a sampling of every sixth 40 μ m thick coronal section (240 μ m) along the rostro-caudal axis of the DG. Cell numbers were expressed as the number of cells per mm³ of GCL. For the SVZ, whose volume directly depends on the size of the proliferating population, a sampling of every sixth 40 μ m thick coronal section (240 μ m) from the most rostral

crossing of the corpus callosum to the start of the third ventricle (crossing of the anterior commissure) (i.e., 1.10 mm to 0.14 mm from bregma) was taken and used for cell quantifications, as described previously [11]. The number of Ki-67⁺ cells per SVZ was then obtained by multiplying the results by six to provide an accurate estimation of the number of cells per SVZ, as previously performed [12–14].

Quantification of Double and Triple Labeled Cells

Analysis of double and triple labeled cells was completed using methods similar to Steiner et al. [15]. Briefly, 1-in-12 series of sections of each animal were double or triple labeled as described above. At least 200 randomly chosen Ki-67⁺ cells within the SGZ or SVZ were analyzed for their coexpression of Ki-67 and markers for the different subpopulations of precursor cells (Fig. 4A and Supporting Information Fig. 4) using a confocal microscope. The percentage of Ki-67⁺ that were positive for the marker was then multiplied by the number of proliferating Ki-67⁺ cells of the corresponding animal previously established as described above. This calculation yielded an accurate estimation of the number of proliferating (i.e., Ki-67⁺)/marker⁺ cells per mm³ of GCL and Ki-67⁺/marker⁺ per SVZ. For cell cycle exit experiments, at least 200 randomly chosen BrdU⁺ cells within the SGZ/GCL or anterior rostral migratory stream (RMS) were analyzed for their coexpression of Ki-67.

Cell Cycle Kinetics

To calculate the total length of the cell cycle (T_C) and the length of S-phase (T_S), the mean labeling index (LI, representing the proportion of BrdU⁺ cells to the number of proliferating, i.e., Ki-67⁺, cells) was plotted as a function of time after application of BrdU (Fig. 6B). Ki-67 is expressed during all phases of the cell cycle, except G_0 , which makes it an appropriate growth fraction (GF) marker [16, 17]. The GF (representing the proportion of proliferating cells) was determined as the maximum LI value attained for each genotype. T_C and T_S were determined using (a) y-intercept = $GF \times T_S/T_C$ and (b) time to reach maximum LI = $T_C - T_S$ [18]. Mitotic cells were identified using antiphosphohistone H3 antibodies (mitotic marker). The combined length of G_2 plus M phases (T_{G_2+M}) was determined as the time required to label all mitotic (i.e., Ki-67⁺/pH3⁺) cells with BrdU [18]. The length of G_1 phase (T_{G_1}) was computed using the equation $T_{G_1} = T_C - (T_S + T_{G_2+M})$, as shown previously [18].

Western Blotting

See Supporting Information.

Statistical Analysis

All numerical analysis were performed using GraphPad Prism software version four for Macintosh. Statistical analysis was performed using Student's *t* test or two-way analysis of variance (ANOVA) followed by a Bonferroni's post-test, depending on the data that were analyzed (see figure legends for details). Data are presented as mean with standard deviation of mean (Mean \pm SD). All experiments were performed on four to six different animals per genotype. Differences were considered statistically significant at $p < .05$.

RESULTS

Cdk4 and Cdk6 Are Expressed in SGZ and SVZ Precursors

A detailed analysis of the expression pattern of the G_1 regulators Cdk4 and Cdk6 in the neurogenic regions of the adult brain is lacking. Using confocal microscopy, we first showed that both Cdk4 and Cdk6 were expressed in the SGZ (Fig. 1) and SVZ (Supporting Information Fig. 1). We further

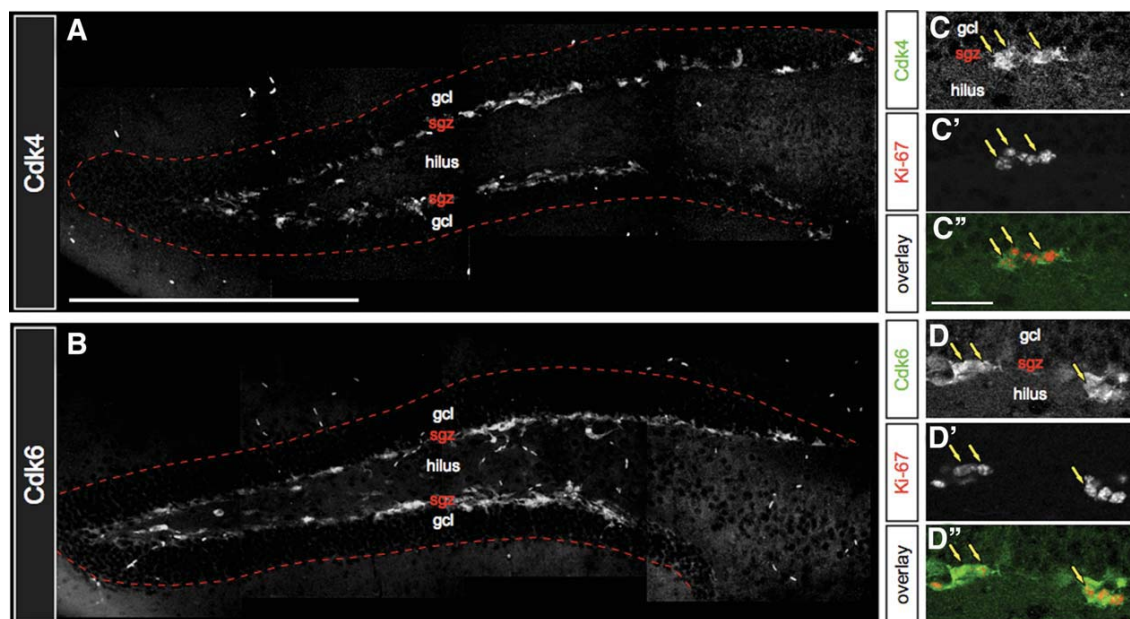


Figure 1. Expression pattern of cyclin-dependent kinase 4 (Cdk4) and Cdk6 in the dentate gyrus (DG). (A, B): Fluorescence images of sections of 6-week-old C57BL/6J DG stained with Cdk4 (A) and Cdk6 antibodies (B). Red dashed lines indicate the outer boundary of the granule cell layer of the DG. Note that the labeling of both Cdks is restricted to the subgranular zone of the DG. (C, D): Fluorescence images of 1.5 μm thick confocal optical sections of DG stained with Cdk4 (C) and Ki-67 (C') as well as Cdk6 (D) and Ki-67 (D') antibodies. Respective overlays are in (C'') and (D''). Yellow arrows point towards double labeled cells. Note the highly similar pattern between both Cdks and Ki-67. Scale bars = 420 μm (A, B), 38 μm (C–D''). Abbreviations: Cdk, cyclin-dependent kinase; gcl: granule cell layer; sgz: subgranular zone.

characterized the expression patterns of Cdk4 and Cdk6 by performing coimmunolabelings with antibodies directed against the proliferative marker Ki-67 (Fig. 1C, 1D). Confocal microscopy analysis revealed that virtually all Ki-67⁺ cells in the SGZ expressed Cdk4 (95% \pm 4.4%; $n = 3$) or Cdk6 (96% \pm 2.8%; $n = 3$), suggesting that the overall majority of precursors express both Cdk4 and Cdk6. In agreement with these findings, we never detected any colabeling of Cdk4/6 with the postmitotic neuronal marker neuronal nuclei (NeuN) (Supporting Information Fig. 2), supporting that Cdk4/6 expression is restricted to proliferating cells in the adult brain.

Cdk6 Is Critical for Neural Precursor Proliferation in the SGZ and SVZ

The expression patterns of Cdk4 and Cdk6 in the SGZ and SVZ prompted us to hypothesize that these kinases could belong to the intrinsic mechanisms that underlie NPC proliferation in these neurogenic niches. As most Cdk4^{-/-} mice survive until adulthood [9, 19] and Cdk6^{-/-} develop normally [10], we took advantage of these genetic models to examine the contribution of Cdk4/6 to adult NPC proliferation. In agreement with previous studies reporting the dwarfism-like phenotype of Cdk4-deficient mice [9, 19], we first observed that Cdk4^{-/-} brains were smaller and lighter than their WT counterparts at every postnatal stage analyzed, suggesting an important role for Cdk4 during brain development (Supporting Information Fig. 3). On the other hand, Cdk6^{-/-} brains were hardly distinguishable from WT brains, although they displayed a small but significant reduced weight starting from postnatal day 30 (P30) (Supporting Information Fig. 3). To determine the importance of Cdk4/6 in the control of cell proliferation in the SGZ and SVZ, we then quantified the number of actively dividing Ki-67⁺ cells in Cdk4^{-/-} and Cdk6^{-/-}

animals. Quantifications first demonstrated that the number of Ki67⁺ cells in the SGZ and SVZ was similar between Cdk4^{-/-} and their respective WT littermates (Fig. 2A–2C and Supporting Information Fig. 4). On the other hand, we found a twofold reduction in the number of Ki-67⁺ cells in the SGZ and SVZ of Cdk6^{-/-} animals (Fig. 2D–2F and Supporting Information Fig. 4). Importantly, a similar proliferation defect was observed in the brain of 10-week-old Cdk6^{-/-} mice (data not shown). Of note, in accordance with previous work showing that the genetic background influences adult NPC proliferation [20], we observed that WT animals arising from Cdk4 and Cdk6 colonies depict different levels of proliferation (Fig. 2C, 2F). Finally, we showed that Cdk4 and Cdk6 proteins were expressed at similar levels in WT mice coming from Cdk4 and Cdk6 colonies, ruling out any putative contribution of the genetic background and relative Cdk4/6 expression levels to the phenotypic discrepancies observed between Cdk4^{-/-} and Cdk6^{-/-} animals (Supporting Information Fig. 5). Overall, these results demonstrate that although Cdk4 and Cdk6 are both expressed by SGZ and SVZ neural precursors, Cdk6 seems to be more essential to promote their proliferation.

To examine whether the defect in proliferation observed in Cdk6^{-/-} neurogenic areas impairs neuronal cell production, the number of cells coexpressing doublecortin (DCX) and NeuN was quantified in the DG of WT and Cdk6^{-/-} mice (Fig. 3A–3E). These cells are young postmitotic neurons arising from proliferating precursors and therefore represent an accurate indicator of the rate of neurogenesis in the adult hippocampus [21]. There was a reduced number of DCX⁺/NeuN⁺ cells (Fig. 3E) in the absence of Cdk6, which is consistent with the decreased levels of precursor cell proliferation (Fig. 2). Importantly, most newly generated DCX⁺/NeuN⁺ neurons undergo a selection process, during which they are

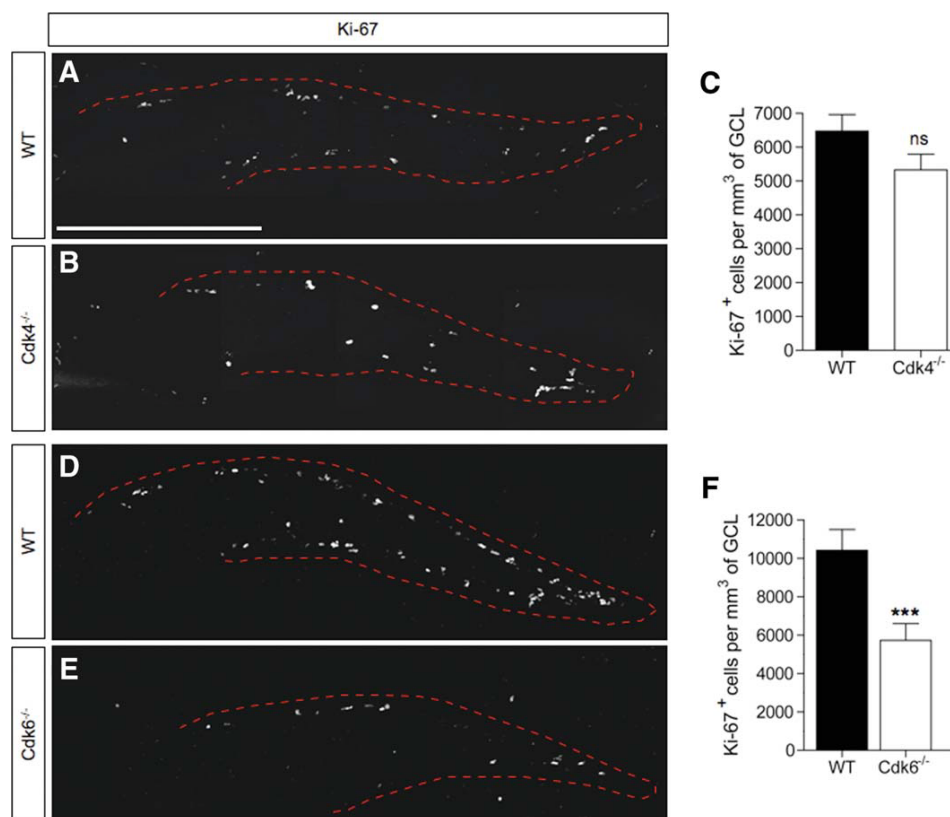


Figure 2. Cell proliferation in cyclin-dependent kinase (Cdk) 4^{-/-} and Cdk6^{-/-} dentate gyrus (DG). (A, B, D, E): Fluorescence images of Ki-67 labeling in the DG of WT (A) and Cdk4^{-/-} (B) as well as WT (D) and Cdk6^{-/-} (E) mice. The red dashed lines delineate the outer boundary of the granule cell layer (GCL). (C, F): Histograms showing the number of Ki-67⁺ cells per mm³ of GCL of the DG in Cdk4^{-/-} (C) and Cdk6^{-/-} (F) brains when compared with their respective aged-matched WT littermates. Scale bar = 400 μ m (A, B, D, E). (C, F): Data are presented as mean \pm SD and were analyzed by a Student's *t* test. *N* is five animals per condition. ***, *p* < .001. Abbreviations: Cdk, cyclin-dependent kinase; GCL, granule cell layer; WT, wild type; ns: not significant.

either recruited into function or eliminated, presumably by apoptosis [22]. Thus, we further analyzed whether Cdk6 deficiency affects the number of newborn neurons surviving in the DG. Hence, we injected BrdU once and quantified the number of BrdU⁺ cells 4 weeks after the injection. We found a significant reduction of the number of surviving BrdU⁺ cells in the Cdk6-deficient DG (Fig. 3J). Triple-labeling experiments of BrdU, NeuN, and astrocytic marker S100 β showed that the vast majority of BrdU⁺ cells became NeuN⁺ neurons 4 weeks after birth regardless of the mouse genotype (Fig. 3F–3I for representative images in WT animals and Fig. 3K). Together with the reduced precursor proliferation and reduced DCX⁺/NeuN⁺ newborn neuron production, these data show that the absence of Cdk6 impedes long-lasting neuronal production in the DG. Similarly, the numbers of BrdU⁺/NeuN⁺ cells were reduced in the OB of mice deficient for Cdk6 4 weeks after the BrdU injection (data not shown). Finally, we sought to determine whether the lack of Cdk6 might affect the survival of newly born cells in the DG. Toward that purpose, we analyzed the expression of activated caspase3 in DG cells and compared the number of BrdU⁺/NeuN⁺ cells observed at 4 weeks to the number of BrdU⁺ cells counted 2 hours after a single BrdU injection to obtain a survival ratio. Neither the number of cells undergoing apoptosis (Fig. 3L) nor the survival ratio (Fig. 3M) were found to be significantly different between WT and Cdk6^{-/-} animals, suggesting that

the net reduction of neuronal production observed in Cdk6-deficient mice does not result from impaired cell survival. Altogether, these data support that Cdk6 is central to adult neurogenesis by regulating the expansion of the precursor pool and hence newborn neuron production.

Cdk6 Selectively Regulates the Proliferation of Neuronally Committed Precursors of the DG

Several types of precursors can be identified in adult neurogenic niches according to their specific morphology and expression of unique sets of molecular markers [23] (Fig. 4A and Supporting Information Fig. 7). Previous works have demonstrated that cell cycle regulation in the adult brain is precursor-type-specific [24], thus we sought to determine whether the defect in proliferation observed in Cdk6^{-/-} brains was restricted to a subpopulation versus all cycling precursors. First, we used glial fibrillary acidic protein (GFAP) and the Sry-related HMG box transcription factor Sox2 to identify immature DG precursors with, respectively, radial (type 1) and nonradial (type 2a) morphology that are not yet restricted to a neuronal fate [25]. No significant differences in the number of proliferating GFAP⁺ (i.e., GFAP⁺/Ki-67⁺) nor Sox2⁺ (i.e., Sox2⁺/Ki-67⁺) precursors were found in Cdk6^{-/-} mice (Fig. 4B–4E). Neuronally committed precursors (types 2b and 3) were identified with antibodies against Neurod1 and DCX

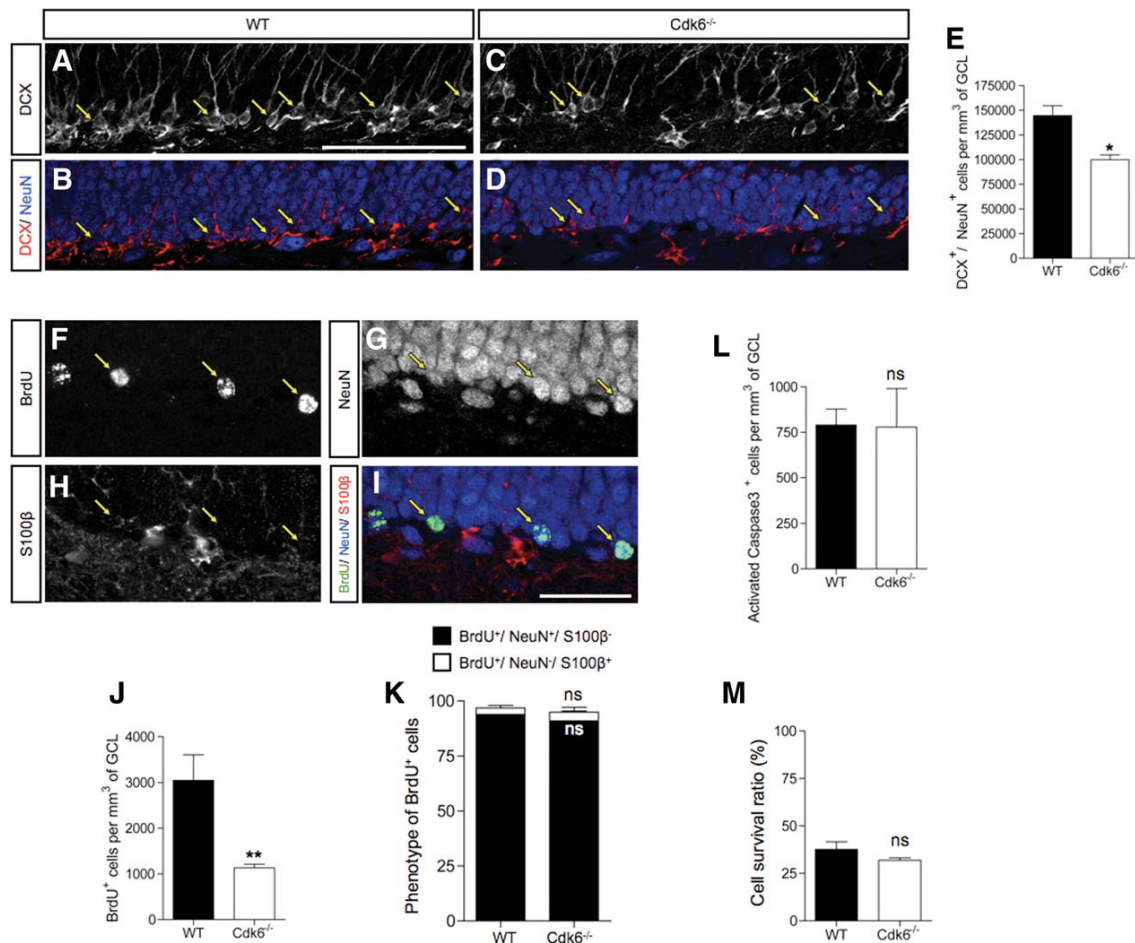


Figure 3. Neurogenesis in cyclin-dependent kinase (Cdk) 6^{-/-} dentate gyrus (DG). (A–D): Fluorescence images of DG stained with doublecortin (DCX) (A, C) and colabeled with NeuN (B, D) in WT (A, B) and Cdk6^{-/-} (C, D) brains. Yellow arrows point toward DCX⁺/NeuN⁺ double-labeled cells. (E): Histograms showing the number of DCX⁺/NeuN⁺ cells per mm³ of granule cell layer (GCL) between WT and Cdk6^{-/-} brains. (F–I): Fluorescence images of 1.5 μm thick confocal optical sections stained with BrdU (F), NeuN (G), S100β (H), and respective overlays (I) 4 weeks after a single BrdU injection in WT animals. Yellow arrows point toward BrdU⁺/NeuN⁺/S100β⁻ cells. (J): Histograms showing the number of BrdU⁺ cells per mm³ of GCL 4 weeks after a single BrdU injection in WT and Cdk6^{-/-} animals. (K): Histograms showing the percentage of BrdU⁺ cells that were either NeuN⁺/S100β⁻ or NeuN⁻/S100β⁺ (black- and white-filled portions of the histograms, respectively) 4 weeks after a single BrdU injection in WT and Cdk6^{-/-} animals. Note that the vast majority of BrdU⁺ became NeuN⁺ (i.e., neurons) within 4 weeks, regardless of the genotype. (L, M): Histograms showing the number of activated caspase3⁺ cells per mm³ of GCL (L) and a BrdU cell survival ratio (M) (formula = #BrdU⁺/NeuN⁺ cells 4 weeks after a single BrdU injection/#BrdU⁺ cells 2 hours after a single BrdU injection) in WT and Cdk6^{-/-} animals. Scale bars = 105 μm (A–D), 42 μm (F–I). (E, J, K, L, M): Data are represented as mean ± SD and were analyzed by a Student's *t* test. *N* is four (E), five (J), five (K), four (L), and five (M) animals per condition. *, *p* < .05; **, *p* < .005. Abbreviations: BrdU, bromodeoxyuridine; Cdk, cyclin-dependent kinase; DCX, doublecortin; GCL, granule cell layer; NeuN, neuronal nuclei; ns, not significant; WT, wild type.

[26]. Quantifications revealed that the numbers of dividing precursors expressing these markers (i.e., Neurod1⁺/Ki-67⁺ and DCX⁺/Ki-67⁺) were all severely lowered in Cdk6^{-/-} mice when compared with their WT littermates (Fig. 4F–4I). Consistently, we also observed that the numbers of neuronally committed precursors expressing Tbr2 (type 2b) [27] and the most committed Prox1-expressing precursor cells (type 3) [15] were reduced in the Cdk6^{-/-} DG (Fig. 4J–4M). Finally, to address whether Cdk6 deficiency affects quiescent precursors of the SGZ, we first quantified the proportion of Sox2⁺ cells that are negative for Ki-67 and detected no difference between WT and Cdk6^{-/-} animals (Supporting Information Fig. 6). To further confirm this result, we tracked quiescent precursors by injecting BrdU twice a day for 1 week followed by a 28-day-chase before sacrifice (Supporting Information Fig. 6). We then counted the number of BrdU label-retaining

quiescent precursors in the DG using BrdU/Sox2/S100β immunohistochemistry. In agreement with our previous data, no significant difference in the number of BrdU⁺/Sox2⁺/S100β⁻ cells was observed between WT and Cdk6^{-/-} mice (Supporting Information Fig. 6). Altogether, these data support that Cdk6 is primarily involved in controlling the proliferation of neuronally committed precursors. We then defined whether Cdk6 deficiency would also preferentially affect the proliferation of a particular subpopulation of precursors of the SVZ. Using Sox2 as a marker of type B cells in the SVZ [28, 29], we first observed that the numbers of Sox2⁺/Ki-67⁺ cells were not impaired in the absence of Cdk6 (Supporting Information Fig. 7). SVZ type B cells continuously give rise to type C transit-amplifying cells themselves producing type A neuroblasts [30, 31] that migrate tangentially along the RMS to the OB, where they differentiate into GABAergic neurons

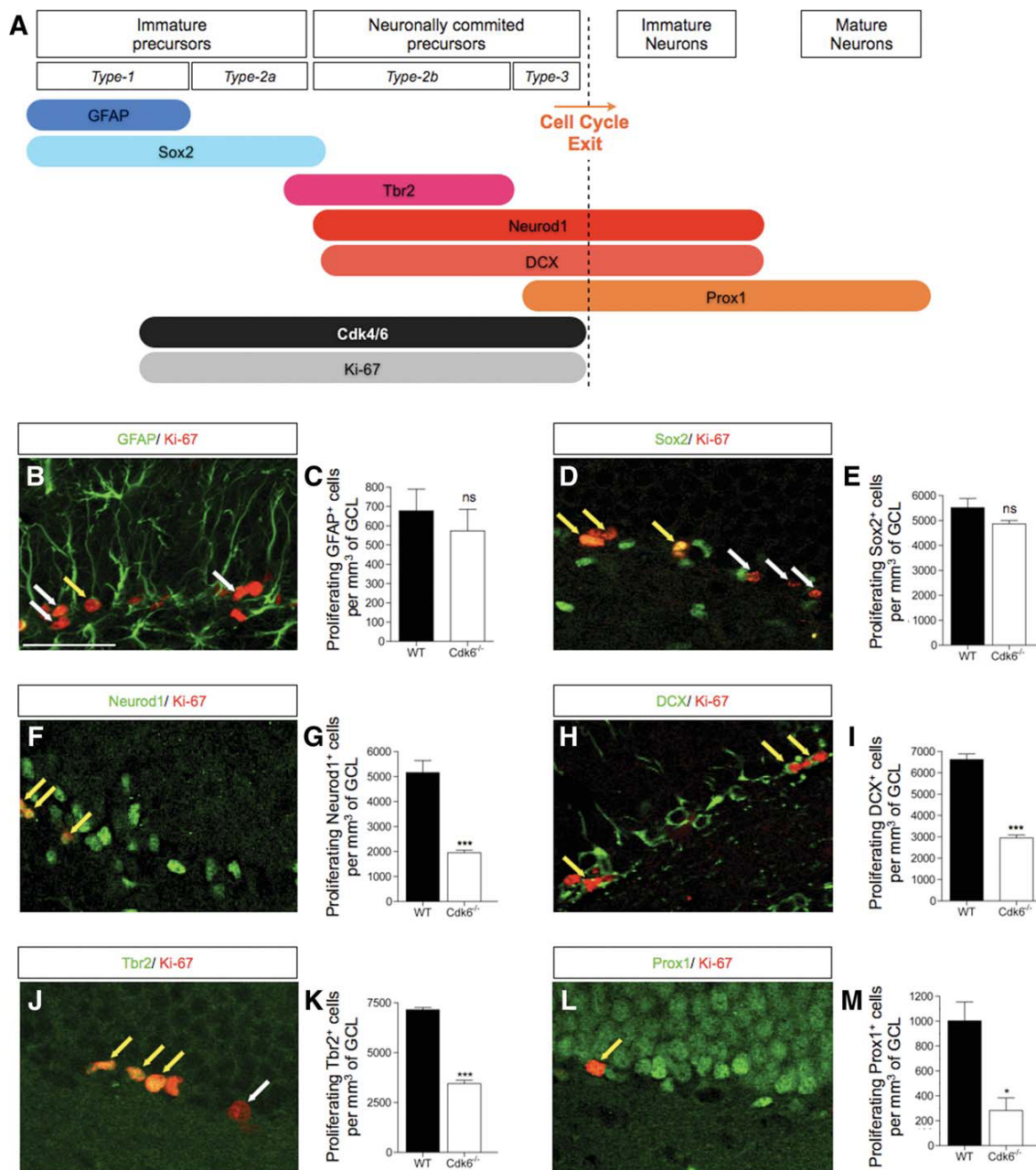


Figure 4. Subtype specific defect in cyclin-dependent kinase (Cdk) 6^{-/-} proliferating dentate gyrus (DG) precursors. (A): Milestones of neuronal development in the adult hippocampus. The expression pattern of Cdk4/6 is also displayed. (B, D, F, H, J, L): Fluorescence images of 1.5 μ m thick confocal optical sections of DG costained with Ki-67 and glial fibrillary acidic protein (GFAP) (B), Ki-67 and Sox2 (D), Ki-67 and Neurod1 (F), Ki-67 and doublecortin (DCX) (H), Ki-67 and Tbr2 (J), and Ki-67 and Prox1 (L) in WT animals. Yellow and white arrows represent double labeled Ki-67⁺/marker⁺ and single labeled Ki-67⁺/marker⁻ cells, respectively. (C, E, G, I, K, M): Histograms showing the number of proliferating (i.e., Ki-67⁺)/GFAP⁺ (C), Ki-67⁺/Sox2⁺ (E), Ki-67⁺/Neurod1⁺ (G), Ki-67⁺/DCX⁺ (I), Ki-67⁺/Tbr2 (K), and Ki-67⁺/Prox1 (M) cells per mm³ of granule cell layer in WT and Cdk6^{-/-} DG. Scale bar = 45 μ m (B, D, F, H, J, L). (C, E, G, I, K, M): Data are presented as mean \pm SD and were analyzed by a Student's *t* test. *N* is four animals per condition. *, *p* < .05; ***, *p* < .001. Abbreviations: Cdk, cyclin-dependent kinase; DCX, doublecortin; GCL, granule cell layer; GFAP, glial fibrillary acidic protein; WT, wild type.

[32, 33]. To determine how critical Cdk6 is for the proliferation of these neuronally committed precursor populations, we quantified the numbers of Dlx2⁺/DCX⁻/Ki-67⁺ (type C) and Dlx2⁺/DCX⁺/Ki-67⁺ (type A) [34]. Both type C and type A proliferating precursors were reduced in the SVZ of Cdk6^{-/-}

mice (Supporting Information Fig. 7). Finally, we did not observe any defect in the number of BrdU label-retaining quiescent precursors in the SVZ (Supporting Information Fig. 6), suggesting that, similarly to the DG, Cdk6 principally controls the expansion of neuronally committed cells in the SVZ.

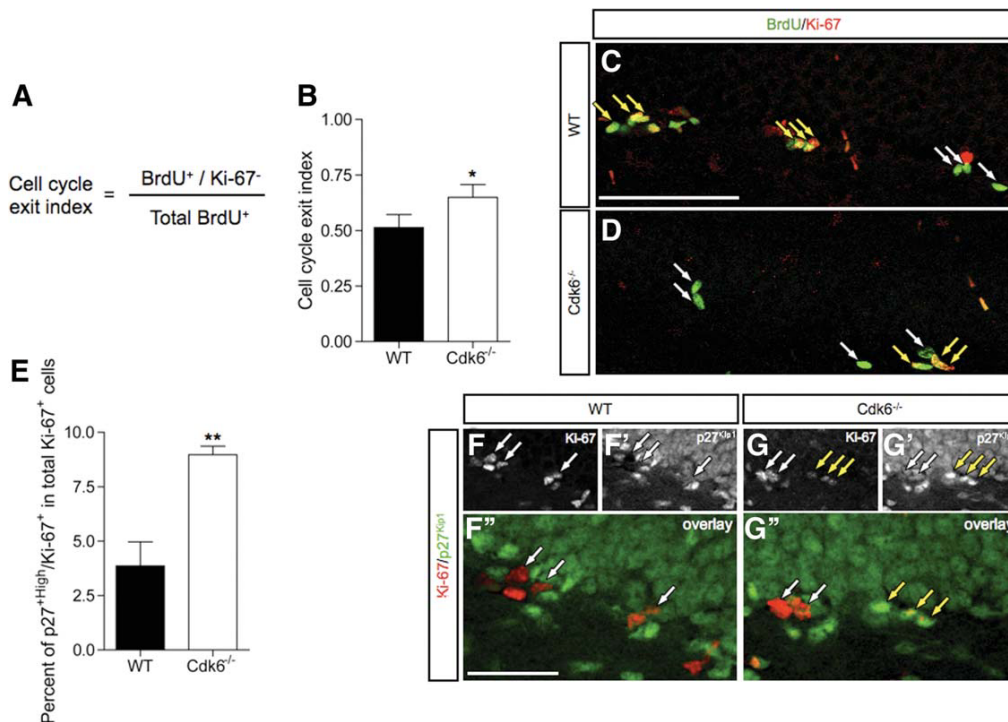


Figure 5. Cell cycle exit in cyclin-dependent kinase (Cdk) $6^{-/-}$ dentate gyrus (DG). (A): Ratio that was used to calculate the cell cycle exit index. (B): Histograms showing the cell cycle exit index in WT and Cdk6 $^{-/-}$ DG precursors. (C, D): Representative images of 1.5 μm thick confocal optical sections of DG costained with Ki-67 and BrdU 24 hours after the injection. Yellow and white arrows in (C, D) represent Ki-67 $^{+}$ /BrdU $^{+}$ (i.e., cells that were incorporated BrdU 24 hours before sacrifice and either remained in the cell cycle or started a new cell cycle) and Ki-67 $^{-}$ /BrdU $^{+}$ (i.e., cells that incorporated BrdU and exited the cell cycle within the following 24 hours), respectively. (E): Histograms showing the percentage of p27 $^{\text{Kip1+High}}$ /Ki-67 $^{+}$ in the total Ki-67 $^{+}$ cells in WT versus Cdk6 $^{-/-}$ subgranular zone. (F–G''): Fluorescence images of 1.5 μm thick confocal optical sections of DG costained with Ki-67 (F, G) and p27 $^{\text{Kip1}}$ (F', G') and respective overlays (F'', G'') in WT (F–F'') and Cdk6 $^{-/-}$ animals (G–G''). White and yellow arrows in (F–F''), (G–G'') represent Ki-67 $^{+}$ /p27 $^{\text{Kip1+Low}}$ and Ki-67 $^{+}$ /p27 $^{\text{Kip1+High}}$ cells, respectively. Scale bars = 70 μm (C, D), 50 μm (F'', G''), 100 μm (F–F'', G–G''). Data are presented as mean \pm SD and were analyzed by a Student's t test. (B, E): N is five (B) and four (E) animals per condition. *, $p < .05$; **, $p < .005$. Abbreviations: BrdU, bromodeoxyuridine; Cdk, cyclin-dependent kinase; WT, wild type.

Cdk6 $^{-/-}$ Neural Precursors Prematurely Exit the Cell Cycle and Display a Lengthened G $_1$ Phase

It is currently assumed that neuronally committed precursors undergo a defined number of cell divisions before exiting the cell cycle and differentiating into neurons [26]. As these precursors give birth to postmitotic neurons, we hypothesized that the selective reduction of their pool could at least partially result from an enhanced rate of cell cycle exit. Hence, we injected mice with a single dose of BrdU, 24 hours before sacrifice. The cell cycle exit index was then scored by defining the ratio between BrdU $^{+}$ /Ki-67 $^{-}$ cells and total BrdU $^{+}$ cells in the DG, corresponding to the fraction of precursors leaving the cell cycle within 24 hours, as described previously [35–38] (Fig. 5A). The cell cycle exit index was significantly increased in Cdk6 $^{-/-}$ (Fig. 5B–5D) but not Cdk4 $^{-/-}$ (Cdk4WT: 47.88% \pm 6.82%; Cdk4 $^{-/-}$: 51.11% \pm 2.73%; $p = .675$; $n = 4$) mice when compared with their respective age-matched WT littermates. Similarly, the cell cycle exit probability of future OB interneurons was increased in the anterior part of the RMS of Cdk6 $^{-/-}$ animals (Supporting Information Fig. 8). It has been shown that the Cdk inhibitor p27 $^{\text{Kip1}}$ progressively accumulates during G $_1$ phase in precursors as increased numbers of cell divisions occur [38, 39]. Using immunofluorescence, we distinguished two types of

p27 $^{\text{Kip1+}}$ cells in the DG, both of them exhibiting nuclear staining. Some cells predominantly found in the SGZ were strongly positive for p27 $^{\text{Kip1}}$ (i.e., p27 $^{\text{Kip1+High}}$), whereas others scattered throughout the GCL and the hilus of the DG showed weaker stainings of p27 $^{\text{Kip1}}$ (i.e., p27 $^{\text{Kip1+Low}}$) (Fig. 5F', 5G'). Colabeling with NeuN revealed that the p27 $^{\text{Kip1+Low}}$ cells were postmitotic NeuN $^{+}$ neurons (data not shown) as described previously [38]. Based on the findings by Durand et al. [39], we considered that Ki-67 $^{+}$ /p27 $^{\text{Kip1+High}}$ cells were about to exit the cell cycle. As shown in Figure 5E, the proportion of Ki-67 $^{+}$ precursors that were p27 $^{\text{Kip1+High}}$ was significantly increased in Cdk6 $^{-/-}$ when compared with WT (Fig. 5F–5G''), which is consistent with the above-mentioned data (Fig. 5B–5D). Altogether, these results strongly suggest that Cdk6 controls the cell cycle exit of neuronally committed precursors, thereby regulating the production of postmitotic neurons.

Recently, Lange et al. [40] demonstrated that forcing the lengthening of G $_1$ is sufficient to promote the exit of the cell cycle of cortical progenitors and their differentiation into projection neurons. As our data support an increased rate of precursor cell cycle exit in Cdk6 $^{-/-}$ neurogenic regions, we compared the length of G $_1$ in Cdk6-knockout and WT animals. To this aim, we performed cumulative S-phase labeling with

BrdU [18, 40] (Fig. 6A–6D). Using this method, we found that the overall duration of the cell cycle (T_C) in DG cycling precursors was increased in $Cdk6^{-/-}$ (22h32') when compared with WT (19h55') (Fig. 6J). Importantly, the T_C of WT DG cells presented in this study fits those calculated previously [41, 42]. The maximal BrdU mitotic LI was reached 3 hours after the initial BrdU injection, regardless of the genotype (Fig. 6E–6I), indicating that the combined length of G_2+M phases was comparable in WT and $Cdk6^{-/-}$ DG precursors (Fig. 6J). Conversely, G_1 showed a selective lengthening (+28.08%) in $Cdk6^{-/-}$ when compared with WT controls, whereas S-phase duration was shortened (–8.31%) (Fig. 6J). In accordance with the phenotype observed in the DG, SVZ precursors also showed a significant lengthening of G_1 in the absence of $Cdk6$ (+82.21%; supporting Information Fig. 8), suggesting $Cdk6$ as a regulator of G_1 length in both neurogenic niches. Finally, we sought to determine whether this defect was affecting all cycling precursors. Thus, we analyzed BrdU incorporation in immature $Neurod1^-$ and neuronally committed $Neurod1^+$ cycling precursors (i.e., $Neurod1^-/Ki-67^+$ and $Neurod1^+/Ki-67^+$, respectively) 6 hours after the initial BrdU injection. Interestingly, we first observed that the LI of immature $Neurod1^-$ cells was higher when compared with more committed $Neurod1^+$ cells in WT brains (Fig. 6K). Second, although the LI of $Neurod1^-$ cells remained unchanged, we observed a significant decrease of the LI of $Neurod1^+$ cells in the absence of $Cdk6$ (Fig. 6K). These results suggest that the overall defect in proliferation observed in the DG of $Cdk6$ -deficient animals mainly results from the impaired proliferation kinetics of neuronally committed precursors.

During cell cycle progression, $Cdk6$ phosphorylates pRb in G_1 , leading to a release of E2f transcription factors and, subsequently, S-phase entry [43]. Similarly to our results, Malumbres et al. [10] previously demonstrated a lengthening of G_1 in $Cdk6^{-/-}$ T lymphocytes on mitogenic stimulation. This defect was concomitant with a delay of pRb phosphorylation on Ser780, a residue specifically targeted by $Cdk4/6$ [44]. Consequently, we hypothesized that the lengthened G_1 measured in $Cdk6^{-/-}$ neural precursors could partially result from pRb hypophosphorylation. Thus, we microdissected the SVZ of WT and $Cdk6^{-/-}$ mice and carried out western blot analysis. Our results revealed a reduction of the phosphorylation of pRb on Ser780 in $Cdk6^{-/-}$ SVZ precursors (Fig. 6L, 6M). Conversely, in agreement with the absence of proliferation defect in $Cdk4^{-/-}$ brains (Fig. 2), we did not detect any significant difference of pRb phosphorylation on Ser780 in $Cdk4^{-/-}$ SVZ precursors (Fig. 6L, 6N). These observations suggest that: (a) $Cdk4$ (or any other kinase) cannot compensate efficiently for the lack of $Cdk6$ pRb-kinase activity in adult neural precursors; (b) the lengthening of G_1 observed *in vivo* in $Cdk6^{-/-}$ adult neural precursors may arise from a reduced phosphorylation status of pRb.

Seizure-Induced Proliferation and Neurogenesis in $Cdk6^{-/-}$ Mice

Acute seizures have been demonstrated to dramatically induce precursor proliferation in the DG [45–47]. Additionally, a recent report from Varodayan et al. [48] demonstrated that seizure leads to a shortening of the cell cycle in DG precursors, suggesting a link between G_1 phase length and seizure-induced proliferation. To address whether $Cdk6$ plays a role during seizure-induced proliferation, we injected kainic acid (KA) to WT and $Cdk6^{-/-}$ animals and quantified precursor cell proliferation in the DG 72 hours later, with a protocol that was described previously [49, 50]. Of note, we consistently observed a large overlap between $Cdk4/6$ and Ki-67

expression 72 hours after either NaCl or KA administration (Supporting Information Fig. 9). As expected (Fig. 2), we found a decreased number of proliferating Ki-67⁺ cells in NaCl-treated $Cdk6^{-/-}$ when compared with WT brains (Fig. 7A, 7C, 7E). However, our cell countings revealed that the levels of precursor proliferation increased drastically following KA treatment in both genotypes (Fig. 7A–7E). Because seizure has been demonstrated to initially trigger the proliferation of immature/noncommitted precursors [49, 50], we then quantified the numbers of GFAP⁺/Ki-67⁺ and Sox2⁺/Ki-67⁺ cells following KA injection. We observed a dramatic increase of the number of GFAP⁺ and Sox2⁺ proliferating precursors, independently of the mouse genotype (Fig. 7F, 7G). These findings are consistent with our data obtained in basal conditions (Fig. 2) and further confirm that the regulation of immature/noncommitted precursor proliferation and cell cycle kinetics occur in a $Cdk6$ -independent manner. More generally, these data preclude a role for $Cdk6$ during the initial phase of seizure-induced proliferation. In addition to transiently enhancing cell proliferation, seizures have been shown to increase the production of newborn neurons [51]. Importantly, these newborn neurons exhibit dramatic changes of neuronal polarity and migration, and integrate aberrantly into the pre-existing neuronal network [52–54], leading to long-term structural changes in the hippocampal circuitry that might contribute to chronic epilepsy [55, 56]. To evaluate the consequences of $Cdk6$ deficiency on seizure-induced neurogenesis, we injected WT and $Cdk6^{-/-}$ animals with BrdU 72 hours following seizure induction and sacrificed the animals 4 weeks later to quantify the number of BrdU⁺ surviving cells. Consistently with our data obtained in basal conditions, $Cdk6$ -deficiency impaired the production of newborn cells in the epileptic DG (Fig. 7H). Altogether, these data suggest that although KA increases the proliferation of immature/noncommitted $Cdk6^{-/-}$ NPCs to a similar extent as in WT, the absence of $Cdk6$ largely prevents the burst in neuronal production observed in WT conditions.

DISCUSSION

Our findings demonstrate a unique and crucial role for $Cdk6$ in controlling neuronally committed precursor expansion and the rate of neuronal production in the adult brain (Supporting Information Fig. 10). Besides, we identified G_1 phase duration as one key regulator of this process.

During brain development, a progressive increase in rates of neuron production is accompanied by increasing frequencies of differentiative divisions that come together with the extension of G_1 phase length [57, 58]. Particularly, proliferative divisions are characterized by a short G_1 , whereas differentiative or neurogenic divisions exhibit a longer G_1 [57, 58]. These findings led to the “cell cycle length hypothesis,” stating that there is an inverse correlation between G_1 phase duration and the onset of neuronal differentiation [59]. Considering (a) type 1 precursors are mainly slowly dividing and account for a small part of the total proliferative activity of the DG (ref. [60] and the present study) and (b) there is no overlap between the expression of Sox2 and $Neurod1$ in the DG (refs. [61, 62] and our unpublished observations), the lower LI of $Neurod1^+$ cells when compared with $Neurod1^-$ cells suggests that immature actively proliferating precursors (type 2a) cycle more rapidly than do neuronally committed precursors (types 2b and 3), supporting for the first time the “cell cycle length hypothesis” for adult neurogenic niches. Moreover, the lengthening of G_1 observed in the absence of

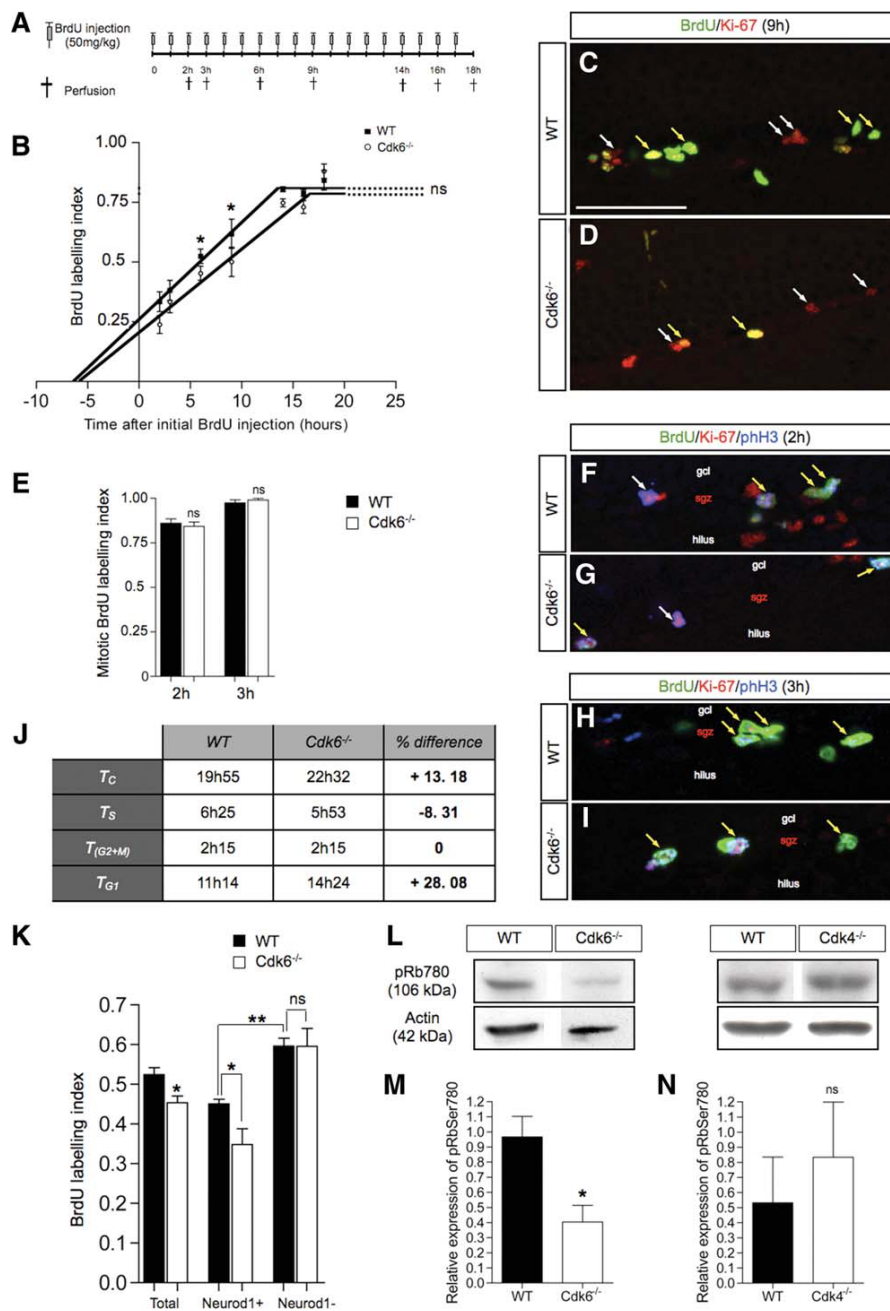


Figure 6. Cell cycle kinetics in cyclin-dependent kinase (Cdk) $6^{-/-}$ dentate gyrus (DG). **(A):** Time course of cumulative BrdU labeling in WT and Cdk6 $^{-/-}$ DG. BrdU was injected every hour (syringes). Mice were sacrificed (crosses) 2, 3, 6, 9, 14, 16, and 18 hours after the first BrdU injection. **(B):** Cumulative BrdU labeling curves of WT (black squares) and Cdk6 $^{-/-}$ (open circles) precursors. Dashed lines indicate growth fractions that were not different between WT and Cdk6 $^{-/-}$ animals. **(C, D):** Representative fluorescence images of 1.5 μ m thick confocal optical sections of DG stained with BrdU and Ki-67 9 hours after the first BrdU injection in WT **(C)** and Cdk6 $^{-/-}$ **(D)** animals. Yellow and white arrows point toward Ki-67 $^{+}$ /BrdU $^{+}$ and Ki-67 $^{+}$ /BrdU $^{-}$ cells, respectively. **(E):** Mitotic BrdU labeling indices of WT and Cdk6 $^{-/-}$ DG after 2 (left) or 3 (right) hours of BrdU exposure. Mitotic labeling index 1.00 = all mitotic (i.e., Ki-67 $^{+}$ /pH3 $^{+}$) cells stained with BrdU. **(F–I):** Representative fluorescence images of 1.5 μ m thick confocal optical sections stained with BrdU, Ki-67, and pH3, 2 **(F, G)** and 3 **(H, I)** hours after the first BrdU injection in WT **(F, H)** and Cdk6 $^{-/-}$ **(G, I)** animals. Yellow and white arrows point toward Ki-67 $^{+}$ /BrdU $^{+}$ /pH3 $^{+}$ and Ki-67 $^{+}$ /BrdU $^{-}$ /pH3 $^{+}$ cells, respectively. **(J):** Table showing the total cell cycle time, S-phase, G2 and M phases, and G1 phase duration in WT and Cdk6 $^{-/-}$ DG. **(K):** Six-hour BrdU labeling indices of WT (black) and Cdk6 $^{-/-}$ (white) precursors among total (left), Neurod1 $^{+}$ (middle), and Neurod1 $^{-}$ (right) precursors. **(L):** Western blot against pRbSer780 in WT and Cdk6 $^{-/-}$ (left), WT and Cdk4 $^{-/-}$ (right) subventricular zone precursors. **(M, N):** Histograms showing the corresponding Western blot quantifications. Relative intensity of pRbSer780 signal was calculated by doing the ratio with the actin signal from the same sample. Scale bars = 65 μ m **(C, D)**, 55 μ m **(F–I)**. Data are the mean of four different animals per genotype per time point \pm SD for the cell cycle kinetics analysis **(B, E, J)** and were analyzed by a two-way ANOVA followed by a Bonferroni's post-test **(K)**. Data are the mean of four different animals of each genotype \pm SD and were analyzed by a Two-way ANOVA followed by a Bonferroni's post-test. **(M, N):** Data are the mean of four proteic extracts coming from four different animals of each genotype \pm SD and were analyzed by a Student's *t* test. * $p < .05$; ** $p < .005$. Abbreviations: BrdU, bromodeoxyuridine; Cdk, cyclin-dependent kinase; gcl: granule cell layer; ns: not significant; sgz: subgranular zone; T_C , total cell cycle time; T_{G2+M} , total G2 and M phases duration, T_{G1} , total G1 phase duration; T_S , total S-phase duration; WT, wild type.

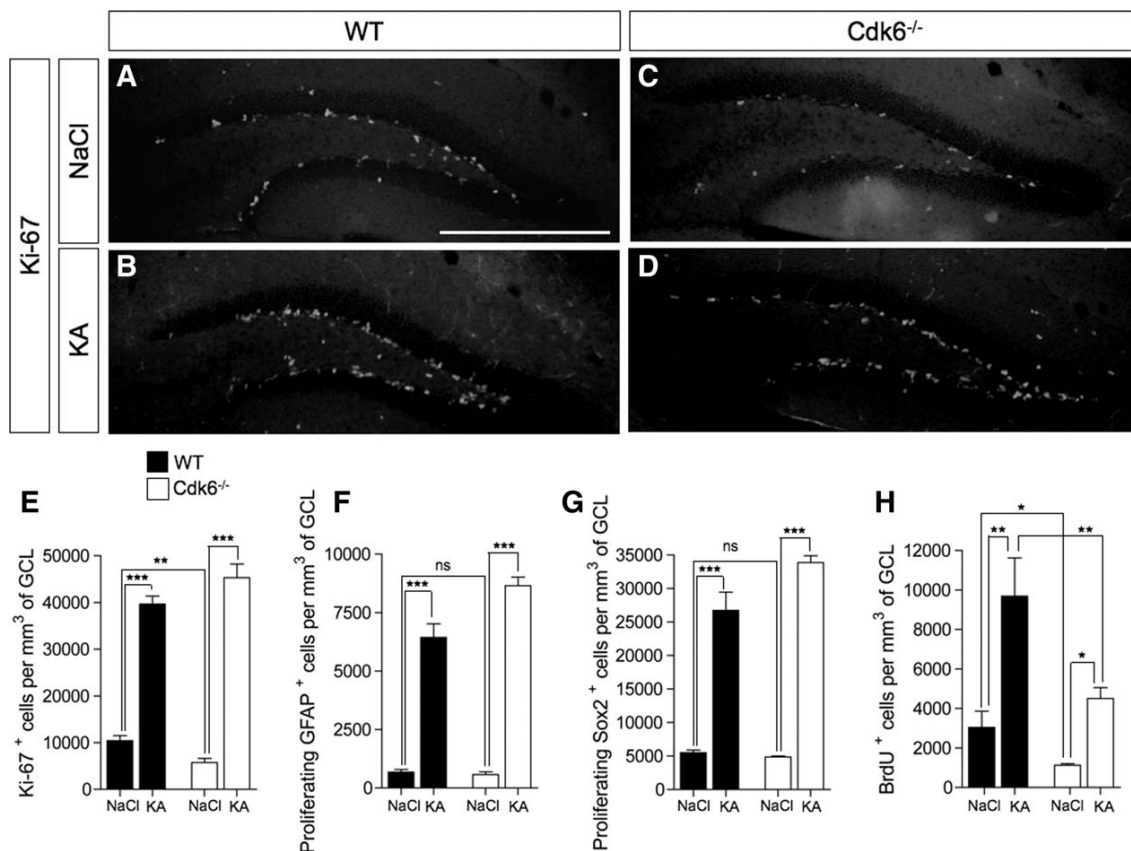


Figure 7. Seizure-induced neurogenesis in wild type (WT) and cyclin-dependent kinase (Cdk) 6^{-/-} animals. (A–D): Representative fluorescence images of Ki-67 labeling in the dentate gyrus (DG) of saline (NaCl)-WT (A) and Cdk6^{-/-} (C) and kainic acid (KA)-treated WT (B) and Cdk6^{-/-} (D) 72 hours after the NaCl or KA injection. (E–H): Histograms showing the number of Ki-67⁺ cells (E), the number of proliferating (i.e., Ki-67⁺/glial fibrillary acidic protein⁺ (F), Ki-67⁺/Sox2⁺ (G) cells, and the number of BrdU⁺ cells 4 weeks after a single BrdU injection (performed 72 hours after the KA injection) (H) per mm³ of granule cell layer in saline- and KA-treated WT and Cdk6^{-/-} DG. Scale bar = 400 μ m (A–D). Data are presented as mean \pm SD. Statistical analysis were two-way ANOVA followed by a Bonferroni's post-test. (E–H): *N* is four different animals per condition. *, *p* < .05; **, *p* < .005; ***, *p* < .001. Abbreviations: BrdU, bromodeoxyuridine; Cdk, cyclin-dependent kinase; GCL, granule cell layer; GFAP, glial fibrillary acidic protein; KA, kainic acid; ns: not significant; WT, wild type.

Cdk6 causes precocious cell cycle exit and differentiation. Interestingly, as the absence of Cdk6 specifically results in the reduction of both the number and the LI of neuronally committed precursors, one could conceptualize a model where Cdk6, presumably through G₁ length regulation, actively promotes proliferative divisions of type 2b cells, preventing them to prematurely give rise to more differentiated type 3 cells and ultimately to neurons. Furthermore, evidences show that type 3 cells can also divide [15] and the absence of Cdk6 could also interfere with their proliferative capacity. Importantly, we did not detect any defect in immature precursors either in basal nor in seizure induced proliferation, suggesting that Cdk6-independent mechanisms efficiently control their proliferation. However, in contrast with WT animals, the dramatic increase of proliferation we observed in Cdk6^{-/-} mice following KA administration did not lead to a proportional increase of newborn neuron production. These findings suggest that a pharmacological inhibition of Cdk6 may be effective in preventing the production of ectopic granule cell neurons, a putative cause of temporal lobe epilepsy in humans [55, 56]. More generally, our results underline the need to decipher the distinct molecular mechanisms controlling G₁ in

the different proliferating populations of the adult brain in normal or pathological conditions.

This work shows that although Cdk4 is expressed in SGZ and SVZ precursors, this kinase is dispensable for cell proliferation in adult neurogenic niches, at least in our experimental conditions. Our results are in accordance with other studies that have previously shown that Cdk4 deficiency does not impede cell proliferation in several highly proliferating tissues, such as skin, small intestine, or hematopoietic system [63, 64]. In addition, Cdk4 is dispensable for liver regeneration following partial hepatectomy [64], an assay considered to be one of the most stringent to evaluate the proliferative capacity of an adult tissue [65]. Altogether, these data support that Cdk4 is, with the exception of certain cell types [9, 19, 66, 67], overall dispensable for adult tissue homeostasis.

Interestingly, we recently demonstrated that another major interphase Cdk, Cdk2, was also dispensable for hippocampal neurogenesis [68]. Importantly, while the long held belief was that all Cdks were expressed in most proliferating cell types [43, 69], we were able to detect Cdk1, Cdk4, and Cdk6 but not Cdk2 expression in the DG and SVZ (our unpublished observations), although a recent article reported a critical role

for the latter in SVZ cell proliferation and self-renewal [70]. Since recent work by Santamaria et al. [71] described a central role for Cdk1 during embryogenesis, it would be interesting to assess if Cdk1 activity also accounts for proliferation in the DG and SVZ as basal levels of both phosphorylation of pRb and cell proliferation are maintained in Cdk6 mutant brains.

Of interest, one major discrepancy between Cdk4 and Cdk6 is the specific role of the latter in blocking differentiation of multiple lineages [69]. For instance, although Cdk4 and Cdk6 are expressed all along the hematopoietic lineage, Cdk6 deficiency specifically impaired expansion of lineage-committed populations rather than proliferation of early hematopoietic precursors [10], a situation mimicking the phenotype we observed in the brain. Interestingly, our results support the idea that the specific involvement of Cdk6 in neural precursors may partially rely on its unique capacity to phosphorylate pRb on Ser780 in these cells. Cdk4^{-/-} embryonic fibroblasts have been previously shown to display reduced levels of pRb-Ser780 while expressing Cdk6 [9], suggesting that the main D-type Cdk responsible for phosphorylation of this residue could be cell-type dependent. Alternatively, it might be conceivable that Cdk4/6 activity itself depends on the cell type [72].

CONCLUSION

Our work unravels a specific function for the G₁ regulator Cdk6 in the control of adult neurogenesis. More generally, this report is, to our knowledge, the first to show that a modification of G₁ duration influences precursor expansion versus differentiation in an adult tissue and therefore contributes to a

better understanding of the mechanisms that underlie the balance between proliferation and differentiation of somatic precursor cells.

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DISCLOSURE OF POTENTIAL CONFLICTS OF INTEREST

The authors indicate no potential conflicts of interest.

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