Development/Plasticity/Repair

# Developmental Roles of p73 in Cajal-Retzius Cells and **Cortical Patterning**

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To examine the role of the p53 homolog p73 in brain development, we studied  $p73^{-/-}$ ,  $p73^{+/-}$ ,  $E2F1^{-/-}$ , and reeler mutant mice. p73 in developing brain is expressed in Cajal-Retzius (CR) cells, the cortical hem, and the choroid plexus. p73-expressing CR cells are lost in  $p73^{-/-}$  embryos, although Reelin is faintly expressed in the marginal zone. Ectopic neurons in the  $p73^{-/-}$  preplate and cortical hem at embryonic day 12 implicate p73 in the early developmental program of the cortex; however, preplate partition and early cortical plate formation are not disturbed. Postnatal  $p73^{-/-}$  mice show a mild hypoplasia of the rostral cortex and a severely disrupted architecture of the posterior telencephalon. In the developing  $p73^{-/-}$  hippocampus, the most striking abnormality is the absence of the hippocampal fissure, suggesting a role of p73 in cortical folding.  $p73^{+/-}$  mice have a less severe cortical phenotype; they display a dorsal shift of the entorhinal cortex and a reduced size of occipital and posterior temporal areas, which acquire entorhinal-like features such as Reelinpositive cells in layer II. CR cells appear unaffected by heterozygosity. We relate the malformations of the posterior pole in p73 mutant mice to alterations of p73 expression in the cortical hem and suggest that p73 forms part of an early signaling network that controls neocortical and archicortical regionalization. In mice deficient for the transcription factor E2F1, a main activator of the TAp73 (transactivating p73) isoform, we find a defect of the caudal cortical architecture resembling the p73<sup>+/-</sup> phenotype along with reduced TAp73 protein levels and propose that an E2F1-TAp73 dependent pathway is involved in cortical patterning.

Key words: cortical hem; hippocampus; entorhinal cortex; E2F1; TAp73; Reelin

### Introduction

p73 is a homolog of the tumor-suppressor gene p53 (Kaghad et al., 1997). In the brain, it is expressed in Cajal-Retzius (CR) cells, the ependyma, and the choroid plexus (Yang et al., 2000; Meyer et al., 2002). The most striking brain abnormalities of p73 knockout mice are loss of CR cells, hydrocephalus, and dysgenesis of the dentate gyrus (Yang et al., 2000). Neurons in various p73<sup>-/-</sup> brain structures undergo extensive postnatal apoptosis (Pozniak et al., 2002).

p73 is a complex protein that is expressed as transactivating (TA) forms that induce growth arrest and apoptosis and as DeltaN (DNp73) isoforms that lack the transactivating N terminus and are able to inhibit TAp73 and p53 transactivation activities (Yang et al., 2000; Grob et al., 2001; Kartasheva et al., 2002). Expression of these isoforms is controlled by alternative promoters (Jost et al., 1997; Yang et al., 2000). DNp73 has antiapoptotic functions (Pozniak et al., 2000) and is expressed in CR cells,

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whereas TAp73 is present only in a small percentage of p73positive cells (Yang and McKeon, 2000; Yang et al., 2000), raising the question of whether DNp73 and TAp73 play different roles in cortical development. Additional diversity is added by differentially spliced C-terminal variants, of which p73 $\alpha$  is the predominant form in the brain (Yang et al., 2000; Pozniak et al., 2002).

CR cells are early-born neurons in the cortical marginal zone that secrete high levels of Reelin, an extracellular matrix protein necessary for radial migration (D'Arcangelo et al., 1995). It has been proposed that Reelin is required for the process of "preplate partition," when the preplate is split into marginal zone and subplate by the developing cortical plate (Marin-Padilla, 1998; Tissir and Goffinet, 2003). This step is disrupted in reeler mice, in which undivided preplate derivatives form the "superplate" on top of an inverted cortical plate (Caviness 1976, 1982). Because CR cells are the main source of Reelin in the embryonic cortex, we ask here whether their absence disturbs preplate partition and the subsequent architectonic development of the p73<sup>-/-</sup> neocortex in a reeler-like manner.

We reported previously that an important subset of CR cells derive from the cortical hem (Meyer et al., 2002). In the p73deficient mice, we further examine the possible relationships between CR cells, p73 expression, and the cortical hem, which is known as a signaling center important for archicortical and neocortical development (Grove et al., 1998; Grove and Fukuchi-Shimogori, 2003).

In the course of this study, we observed substantial abnormalities of cortical development also in  $p73^{+/-}$  mice. In an attempt to correlate alterations of p73 expression with specific brain defects, we compared  $p73^{+/-}$  and E2F1 knock-out mice. E2F1 belongs to a family of transcription factors that transactivate cell-cycle regulators and proapoptotic genes (Field et al., 1996); it is one of the main activators of the TAp73 promoter (Irwin et al., 2000; Stiewe and Putzer, 2000; Zaika et al., 2001). We report here that E2F1 homozygous and p73 heterozygous mice present similar defects of cortical patterning and that TAp73 protein levels are significantly reduced in the  $E2F1^{-/-}$  brain. Our findings suggest that an E2F1–TAp73-dependent pathway regulates important aspects of cortical development.

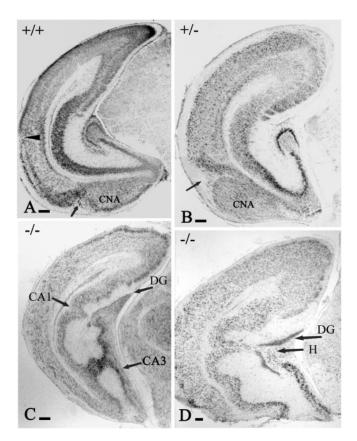
## **Materials and Methods**

We examined  $p73^{-/-}$  and  $p73^{+/-}$  mutant mice and wild-type (WT) littermates (Yang et al., 2000). A total of 63 animals from heterozygous intercrosses (39 prenatal and 26 postnatal animals) were studied at embryonic day 10 (E10), E11, E12, E13, E15, E15.5, and E18, postnatal day 2 (P2), P3, P4, P7, P10, and P12, and adult. Of the 39 embryos, 13 were p73 homozygotes, 15 were heterozygotes, and 11 were WT; genotyping was performed as by Yang et al. (2000). In addition, we studied C57BL/6 mouse brains at various prenatal (E13, E14, E16, and E18) and postnatal ages. The E10-E15.5 embryos were fixed in 4% paraformaldehyde, postfixed in Bouin's fixative, and embedded in paraffin according to standard procedures. Sections were cut serially at 7  $\mu$ m and processed for immunohistochemistry (IHC) and in situ hybridization (ISH). The E18 and postnatal p73 mutant mice were frozen and stored at −80°C. Seven of these brains were cut into 20-um-thick serial sections on a cryostat and also stored at -80°C. For IHC, the frozen sections were air dried, fixed in Bouin's fixative, washed, and then processed as described previously (Meyer et al., 2002). The remaining 19 brains were removed from the freezer and quickly immersed in Bouin's fixative. After fixation for 2 d, they were embedded in paraffin, cut at 10  $\mu$ m, and processed for IHC. ISH was done as described previously (Yang et al., 2000; Meyer et al., 2002). reeler-Orleans (*Reln<sup>rl-Orl</sup>*) mice were studied at E16 and P20. In these mice, a C-terminal truncated protein is produced but not secreted and can be detected using antibodies (Abs) G10 and 142 (de Bergeyck et al., 1997, 1998).  $E2F1^{-/-}$  mice were obtained from The Jackson Laboratory (Bar Harbor, ME). We studied P1, P2, and adult animals. Reeler and E2F1<sup>-/-</sup> mice were fixed in Bouin's fixative, embedded in paraffin, and cut at  $10 \mu m$ .

For IHC, sections were incubated in the primary antibodies overnight in a humid chamber. After rinsing, they were incubated in the corresponding biotinylated secondary antibodies (rabbit anti-mouse IgG or goat anti-rabbit IgG; Dako, Glostrup, Denmark), diluted at 1:200 in Tris-buffered saline (TBS), washed, and incubated in ABC (Dako). Bound peroxidase was revealed using 0.05% 3,3'-diaminobenzidine (Sigma, St. Louis, MO), 0.04% ammonium nickel (II) sulfate, and 0.03% hydrogen peroxide in TBS, pH 7.6. The sections were dehydrated, cleared, and covered with Eukitt (O. Kindler, Freiburg, Germany).

The following primary antibodies were used: rabbit polyclonal anti-p73 antibody (1:150) (Kaghad et al., 1997), mouse monoclonal anti-p73 Ab-2 (1:150) (NeoMarkers, Fremont, CA), mouse monoclonal anti-Reelin antibodies G10 and 142 (1:400) (de Bergeyck et al., 1998), rabbit polyclonal antibody against calretinin (1:2500) (Swant, Bellinzona, Switzerland), and mouse monoclonal antibody against GFAP (1: 250) (NeoMarkers).

Immunoblotting. For total protein extraction, the posterior telencephalons of  $E2F1^{-/-}$  and wild-type brains at P2 were homogenized with lysis buffer (62.5 mM Tris-HCl, 1% SDS, and 10% glycerol, pH 6.8). Protein quantification was done by bicinchoninic acid assay (Smith et al., 1985). Five percent mercaptoethanol and 0.001% bromophenol blue were then added, and samples were boiled at 95°C for 5 min. Equal amounts (40 pg) of each sample were electrophoresed on 10% SDS-PAGE and transferred to Hybond-P membranes (Amersham Biosciences, Little Chalfont, Buckinghamshire, UK). Membranes were preincubated with 5%



**Figure 1.** The  $p73^{+/-}$  and  $p73^{-/-}$  brain phenotypes. A, P3 WT mice. B, P3 +/- mice. C, P3 -/- mice. D, P12 -/- mice, Nissl stain. B, The  $p73^{+/-}$  brain displays abnormal shape and cytoarchitecture of caudal cortical areas and posterior cortical nucleus of the amygdala (CNA). A, B, Arrows point to the periamigdaloid fissure, which is shifted dorsally; the arrowhead in A points to the rhinal fissure, which is unrecognizable in B. C, D, The variability in the malformations of hippocampal architecture in  $p73^{-/-}$  mice. All of the sections are at comparable levels. DG, Dentate gyrus; H, hilus. Scale bars, 200  $\mu$ m.

blotting-grade blocker nonfat dry milk (Bio-Rad Laboratories, Hercules, CA) in TBS with 0.1% Tween 20 (TBS-T) at room temperature for 1 hr and washed in TBS-T. Membranes were incubated with a rabbit polyclonal anti-p73 antibody raised against amino acids 1–15 of human p73 (p73 Ab-5; NeoMarkers) and with a rabbit polyclonal anti-p73 that detects amino acids 427–636 (Kaghad et al., 1997). Membrane incubations with both antibodies (1:500 in TBS-T with 5% BSA) were performed overnight at 4°C. Antibody labeling was developed by incubation with an HRP-conjugated goat anti-rabbit (1:20,000) and visualized with the ECL plus Western Blotting Detection System kit (Amersham Biosciences). Membranes were washed in TBS-T and reprobed with monoclonal mouse anti-tubulin antibody (T 9026; Sigma) (1:100,000 in TBS-T with 5% BSA) for 1 hr, followed by incubation with an HRP-conjugated goat anti-mouse secondary antibody (1:10,000). Specific bands were visualized with the ECL kit, scanned with the GS-800 Calibrated Densitometer, and analyzed with the image analysis program Quantity One (Bio-Rad). For immunosignal quantification, band intensities were normalized to basal values obtained in wild-type brains. Data were obtained in optic density per square millimeter and calculated as the percentage of immunostaining values in E2F1 knock-out brains relative to control brains.

## Results

# Two distinct brain phenotypes in $p73^{-/-}$ and $p73^{+/-}$ mutant mice

On the basis of cytoarchitecture and Reelin expression, we distinguished two main p73 phenotypes (Fig. 1).  $p73^{-/-}$  mice showed architectonic abnormalities of practically all structures of the limbic telencephalon, including hippocampal formation, amygdala,

retrosplenial, and entorhinal cortex (Fig. 1*C,D*). In  $p73^{+/-}$  mice, the alterations were more subtle and affected mainly the entorhinal cortex, the posterior cortical nucleus of the amygdala, and the caudal neocortex (Fig. 1B). In both  $p73^{-/-}$  and p73<sup>+/-</sup> mice, structures at the posterior pole of the telencephalon were more severely affected than rostral ones. The severe size reduction of the caudal cortex in the  $p73^{-/-}$  brain is evident in the parasagittal section of Figure 2 B. In  $p73^{+/-}$  mice, a similar reduction of the posterior pole was first recognized at E18, when the ventricular/subventricular zone extended farther posteriorly in the WT compared with the  $p73^{+/-}$  brain (Fig. 2*C*–*F*).

## The postnatal $p73^{-/-}$ telencephalon

p73<sup>-/-</sup> brains showed the following distinctive features: absence of Cajal-Retzius cells, generalized cortical hypoplasia and caudal dysplasia, severe hippocampal malformations, and abnormal clusters of Reelin-positive stellate cells in layer II of the entorhinal cortex.

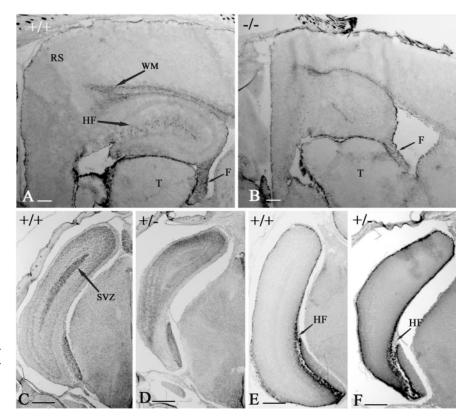
Even in the most severe cases, rostral cortical areas were relatively mildly affected (Fig. 3 *A*, *B*). Lamination and cytoarchitecture were basically preserved, although the cortex was narrower and the white matter and corpus callosum were reduced in size. In contrast, the posterior pole of the hemisphere, including primary and secondary visual areas, the auditory association cortex, and the retrosplenial

cortex, was profoundly altered (Fig. 3*C*). In most cases, these areas had lost the normal lamination pattern and displayed large irregularly arranged cells, many of which were Reelin immunoreactive (IR) (Fig. 3*D*).

The hippocampus presented a complex set of anatomical abnormalities in addition to the absence of the infrapyramidal dentate blade reported by Yang et al. (2000). The hippocampal fissure, which normally separates CA1 and CA2 from the dentate gyrus, was missing. The pyramidal cell layer of CA1 and CA2 was deeply folded, and abnormal cell bridges extended between CA1 and CA3 (Fig. 1*C*) or CA1 and the hilus (Fig. 1*D*).

### p73 in the preplate

In the early preplate (E10–E11), expression of p73 in the WT telencephalon was sparse and mostly close to the roof plate (data not shown). Intense p73 positivity appeared at E12–E12.5 throughout the cortical hem and in CR cells (Meyer et al., 2002). At later embryonic stages, p73 disappeared from the hem and was confined to CR cells and the choroid plexus. ISH, using probes directed against exon 3 (under the control of the first promoter and specific for TAp73 expression) and for exon 3bis (controlled by the second promoter and specific for DNp73 expression), revealed a differential distribution of the respective transcripts. In E12 WT brains, transcripts of exon 3 were restricted to the cortical hem (Fig. 4A, arrowhead), whereas exon 3bis was expressed in the cortical hem and in CR cells all along the cortical wall (Fig. 4D). In E12 p73<sup>-/-</sup> mice, exon 3 transcripts were still present in



**Figure 2.** The posterior pole of the telencephalon. A, P7 WT mice. B, P7 — / — mice, parasagittal sections immunostained with GFAP. Cortical areas at the posterior pole are reduced in size in the  $p73^{-/-}$  mouse. The white matter (WM) area is also smaller, and the hippocampal fissure (HF) is missing. F, Fimbria; RS, retrosplenial cortex; T, thalamus. C–F, The posterior pole of p73 heterozygous (D, F) and WT (C, E) mice at E18 in coronal Nissl (C, D) and Reelin (E, F) stained sections. Sections are at the same level to show the size reduction of the caudal telencephalon and the dorsal shift of entorhinal cortex. The proliferative subventricular zone (SVZ) is still visible in C (WT) but not in D  $(D, F73^{+/-})$ . E, F, The number and distribution of Reelin-IR CR cells in the hippocampal fissure and in the cortical marginal zone are similar in WT and D D D D mice. Scale bars, 200 D m.

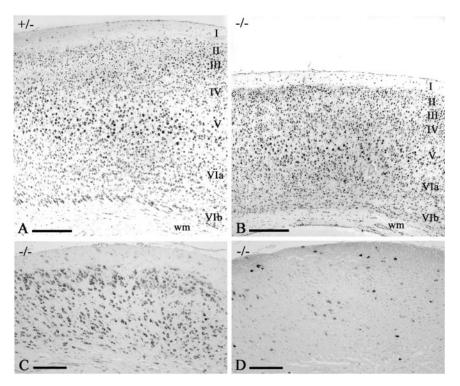
the cortical hem (Fig. 4B, arrowhead), but exon 3bis transcripts were absent (data not shown), indicating that there was no functional TAp73 protein to activate the DNp73 promoter. These findings demonstrate a difference in the expression of p73 in the hem and in CR cells and show that CR cell survival depends on DNp73 expression because they are absent in the  $p73^{-/-}$  brain.

The earliest morphological differences between WT and  $p73^{-/-}$  brains were detected at E12. In the mutant brains, ectopic calretinin-IR cells appeared in the hem and the posteromedial preplate (Fig. 4G), whereas there were no such cells in WT mice of this age (Fig. 4F). Calretinin-IR pioneer cells in the lateral preplate and basal forebrain were also more numerous in knock-outs than in WT mice (Fig. 4F, G), indicating that the loss of p73 gene products changed the developmental program of the cortical neuroepithelium at the preplate stage.

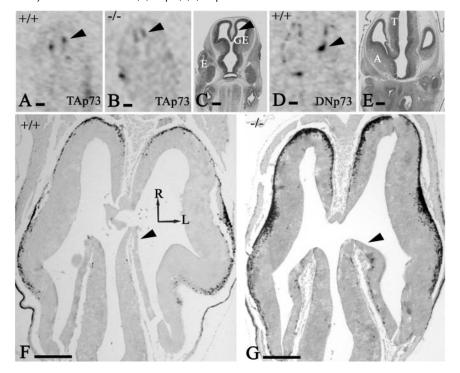
# The development of the cortical plate in the absence of CR cells

We asked whether the absence of CR cells in the E12  $p73^{-/-}$  preplate disrupts preplate partition and early cortical plate development, leading to a reeler-like phenotype. At E15, in WT and  $p73^{+/-}$  embryos, CR cells were intensely IR for Reelin (Fig. 5*A*), p73 (Fig. 5*C*), and calretinin (Fig. 5*E*). Cells with these characteristics were not found in the  $p73^{-/-}$  marginal zone, which contained only few faintly Reelin-IR cells (Fig. 5*B*).

Figure 5*F* shows that, despite the absence of the CR cells, the  $p73^{-/-}$  preplate split into marginal zone and subplate. Calretinin



**Figure 3.** The cortical phenotype of  $p73^{-/-}$  mice. A, B, P12 mice, NissI stain. The frontal (motor) cortex of  $p73^{+/-}$  mice (A) displays normal cytoarchitecture. In the  $p73^{-/-}$  motor cortex (B), basic lamination and cytoarchitecture are preserved, but the cortex is smaller and white matter (wm) is reduced. C, D,  $P73^{-/-}$  (same case as in B), stained with NissI (C) and Reelin (D). In occipital cortex, lamination is lost and cell density is reduced. Irregular clusters of Reelin-IR neurons are present in the profoundly altered cortex. Scale bars: A, B, 200  $\mu$ m; C, D, 100  $\mu$ m.



**Figure 4.** The preplate stage. A, B, D, E12 mice, ISH using specific probes for exon 3 (TAp73) in WT (A), exon 3 in  $p73^{-/-}$  (B), and exon 3bis (DNp73) in WT (D) mice. TAp73 transcripts are expressed in the cortical hem of WT and  $p73^{-/-}$  mice, whereas DNp73 transcripts are present in the WT cortical hem and in CR cells of the cortical preplate. C, E, NissI-stained sections of a control brain at corresponding levels; C for A and B and E for D. The cortical hem is indicated by arrowheads. E, E, E, E12 mice, calretinin IHC, horizontal sections. E, WT mice. E, E, E methods. The mutant preplate presents increased numbers of calretinin-IR pioneer neurons compared with the WT preplate. In addition, there are ectopic calretinin-IR neurons in the cortical hem (arrowheads) that are absent in the WT embryo. Arrows indicate the orientation; E, rostral; E, lateral; E, amygdala; E, eye; E, ganglionic eminence; E, thalamus. Scale bars: E, 200 E, E, 100 E, 10

stains pioneer cells above and below the WT cortical plate (Meyer et al., 1998, 2000); these cells were in their correct position in the  $p73^{+/-}$  and  $p73^{-/-}$  mice (Fig. 5*E*,*F*). Early thalamocortical fibers were also calretinin IR and terminated in their correct target layer, the subplate (Fig. 5 E, F) (Allendoerfer and Shatz, 1994); in contrast, in reeler embryos they traversed the cortical plate to reach the superplate (Fig. 5*G*). The difference between reeler and  $p73^{-/-}$  mice is important insofar as reeler CR cells express p73 but no functional Reelin (Fig. 5D), which leads to a disruption of preplate partition. In  $p73^{-/-}$  mice, very low amounts of Reelin in the marginal zone were able to initiate preplate partition and cortical plate migration; the absence of p73 seemed to play no role in these events.

# The absence of the hippocampal fissure in the $p73^{-/-}$ brain

We compared the development of the hippocampal fissure in E15 and E16 WT,  $p73^{-/-}$ , and reeler embryos. In WT mice, the prospective fissure was marked by an indentation of the pial surface together with an approximately triangular collection of CR cells that coexpressed calretinin, Reelin, and p73 (Fig. 6A–C). The dentate primordium was just ventral to the incipient fissure and characterized by calretinin (Fig. 6A) and proliferating cell nuclear antigen immunoreactivity (data not shown). In  $p73^{-/-}$  mice, CR cells were missing, and the pial surface did not invaginate (Fig. 6D, E). In contrast, in E16 reeler mice, p73/Reelin-IR CR cells were collecting in the fissure, which showed normal folding (Fig. 6F). In fact, the development of the hippocampal fissure is not disturbed in reeler mice.

Examination of the postnatal hippocampus provided additional clues to understand the fissure defect. In the early postnatal WT and  $p73^{+/-}$  mice, a band of GFAP-IR astrocytes extended all along the pial surface into the hippocampal fissure (Fig. 7E), whereas the underlying molecular layers of dentate gyrus and Ammon's horn were densely populated by CR cells (Fig. 7A, C). In  $p73^{-1}$ mice, the hippocampal fissure was reduced to a small indentation of the pial surface that did not extend farther into the brain (Fig. 7*F*). Subpial astrocytes were restricted to the brain surface, and CR cells were missing (Fig. 7*B*,*D*). In consequence, the molecular layers of the dentate and CA1/subiculum were fused instead of being separated by the fissure elements. At approximately P10 in the WT mice, GFAP-IR astrocytes invaded all strata of Ammon's horn, and the radial glia scaffold was established in the dentate (Fig. 7G). In contrast, astrocytes were sparse

and dispersed in  $p73^{-/-}$  mice, and the radial glia scaffold of the dentate gyrus was unrecognizable (Fig. 7H). calretinin-positive cells of the hilus had an abnormal position (Fig. 7C,D), indicating that p73 deficiency severely alters the architecture of all components of the hippocampal formation.

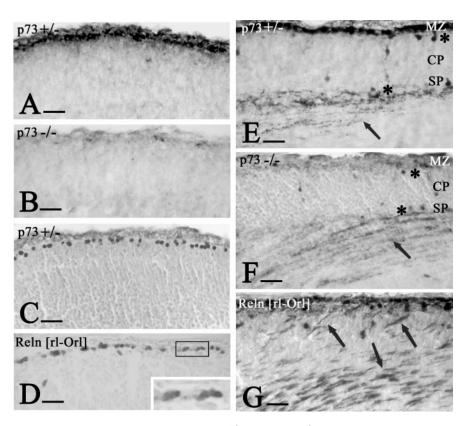
The  $p73^{+/-}$  telencephalon In  $p73^{+/-}$  mice, the cortex appeared mostly normal, except for the most caudal regions, and, most strikingly, the entorhinal cortex. The large projection cells of entorhinal layer II express Reelin (Drakew et al., 1998; Perez-Garcia et al., 2001) and define this area. In  $p73^{+/-}$  mice, the entorhinal cortex was shifted into a more dorsal position, concurrent with an enlargement of the posterior cortical nucleus of the amygdala (Fig. 8A-C). The caudal neocortex (primary and secondary visual cortex and auditory association area) was significantly reduced in size, and the cytoarchitecture appeared disorganized. Furthermore, a band of large Reelin-IR neurons extended from the displaced entorhinal layer II into the altered neocortical territory (Fig. 8A-D), suggesting a transformation of lateral and dorsal neocortex into an entorhinal-like cortex. The presence of ectopic Reelin-IR cells and fibers in the white matter of the posterior pole suggested abnormal axonal projections. Early postnatally, the molecular layer of the altered cortex was populated by numer-

ous CR cells (Fig. 8E,F), which were intensely IR for Reelin (Fig. 8E) and p73 (Fig. 8F). At P12, most CR cells had disappeared (Fig. 8B), as had their counterparts in WT brains.

# The $E2F1^{-/-}$ brain closely resembles the $p73^{+/-}$ phenotype

The transcription factor E2F1 is one of the main activators of TAp73 (Irwin et al., 2000; Stiewe and Putzer, 2000). In early postnatal  $E2F1^{-/-}$  mice, we detected a brain phenotype similar to that of  $p73^{+/-}$  mice, characterized by an increase of the posterior cortical nucleus of the amygdala, a dorsal shift of the entorhinal cortex, and a reduction and entorhinal-like transformation of the caudal neocortex, along with the presence of Reelin-positive cells in layer II of the altered territory (Fig. 9B–D). The rhinal fissure was unrecognizable. Calretinin immunostaining revealed that the laminar distribution of calretinin-IR interneurons in the auditory and visual areas of E2F1<sup>-/-</sup> animals was clearly distinct from the WT pattern (Fig. 9A, B), indicating an alteration of the normal cortical architecture. The white matter of the mutant posterior pole was reduced in size and populated by increased numbers of calretinin- and Reelin-IR interstitial neurons. These anatomical abnormalities developed in the presence of CR cells, which were indistinguishable from those of WT mice. In particular, p73 immunoreactivity in CR cells was apparently unchanged (data not shown) and did not reflect the minor reduction of DNp73 levels observed in the  $E2F1^{-/-}$  brains (see below).

Although this malformation closely resembled the p73<sup>+/-</sup>



**Figure 5.** Early development of the cortical plate. A, E15  $p73^{+/-}$  mice. B, E15  $p73^{-/-}$  mice, Reelin. A, CR cells form a dense cell layer below the pial surface. B, Only few faintly Reelin-positive cells are in the marginal zone (MZ). C, E15  $p73^{+/-}$  mice, high expression of p73 in CR cells. E, E15 p73  $^{+/-}$  mice. F, p73  $^{-/-}$  mice, calretinin. E, CR cells at this stage are calretinin positive and are absent in the p73 knock-out brain. Asterisks mark calretinin-IR pioneer cells above and below the cortical plate (CP), and arrows point to thalamocortical fibers terminating in the subplate (SP); both elements are unchanged in the  $p73^{-/-}$  brain. D, G, E16  $Reln^{rl-Orl}$  mice. D, Reelin and p73 are coexpressed in CR cells. Inset, p73 in the nucleus and Reelin in the cytoplasm of the CR cells. G, Calretinin stains superplate elements and aberrant thalamocortical fibers (arrows) traversing the abnormal CP. Compare with F, in which CP and thalamocortical fibers have normal position. Scale bars, 25  $\mu$ m.

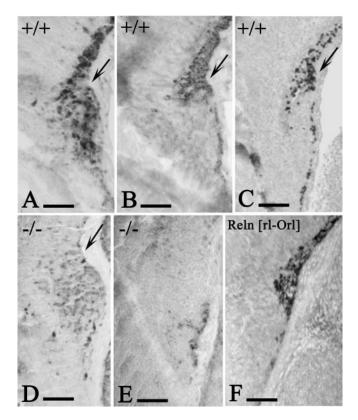
phenotype, the dorsalization of the entorhinal cortex and the reduction of the neocortical territory were less pronounced.

## TAp73 expression is reduced in the $E2F1^{-/-}$ brain

Immunoblot analysis of E2F1-deficient and WT brains at P2 revealed a reduction of TAp73 expression in the mutant (Fig. 10). In WT brains probed with an N-terminal antibody (Fig. 10 A), we observed a large band of 82 kDa corresponding to TAp73 $\alpha$  and a smaller band of 72 kDa corresponding to TAp73β (Yang and McKeon, 2000). In the  $E2F1^{-/-}$  brains, we observed a 50% reduction of the 82 kDa band and a 20% reduction of the 72 kDa band. Immunoblotting using the C-terminal antibody (this antibody consistently stains DNp73-expressing CR cells) revealed a single band of  $\sim$ 66 kDa (Fig. 10C), which is in accordance with the predicted molecular weight of DNp73 $\alpha$  (Yang and McKeon, 2000). This band showed only a 7% reduction in the E2F1 mutant brains. Using the N-terminal antibody, we observed an additional band of higher molecular weight in the E2F1<sup>-/-</sup> brains (Fig. 10A), which could not be attributed to the known p73 isoforms. Our results show that the loss of function of E2F1 leads to a significant reduction of TAp73 protein levels and a minor decrease of DNp73 levels in the brain.

### Discussion

Our study of p73-deficient brains points to extensive activities of p73 in cell proliferation, cell survival, and cortical patterning. The



**Figure 6.** Early development of the hippocampal fissure. A-C, E15 WT mice. D, E, E15  $p73^{-/-}$  mice. F, E16  $Reln^{fl-Orl}$  mice. CR cells express calretinin (A), Reelin (B), and p73 (C) and occupy the fissure (arrows), which begins to fold. In E15  $p73^{-/-}$  embryos (D, calretinin; E, Reelin), CR cells are absent and the fissure does not fold. F, Reelin. In E16  $Reln^{fl-Orl}$  mice, the hippocampal fissure develops normally. Scale bars, 100  $\mu$ m.

dramatic defects of p73 deficiency are difficult to reconcile with its rather selective expression in the brain. We propose that p73 may play a role in the signaling network of the cortical hem, which regulates neocortical and archicortical regionalization and the generation of CR cells. We discuss the possible role of p73 in CR cells, the differential expression of the main products of the p73 gene (TAp73 and DNp73) in early brain development, and the involvement of an E2F1–TAp73-dependent signaling pathway in the regionalization of the posterior telencephalon.

### Cajal-Retzius cells in cortical development

CR cells clearly depend on p73 and do not develop in the p73<sup>-/-</sup> brain. CR cells secrete Reelin, which activates a signaling pathway that leads to phosphorylation of the adapter protein Dab1 in cortical plate cells (Howell et al., 1999). In *reeler* mice, cortical layers are inverted and radial architecture is disturbed (Lambert de Rouvroit and Goffinet, 1998). The absence of CR cells as the main source of Reelin would be expected to produce *reeler*-like defects of cortical migration and lamination. However, lack of CR cells in the p73 mutant is compatible with quite normal lamination and cytoarchitecture, perhaps because Reelin is faintly expressed at the critical stages for preplate (Meyer et al., 2002) and cortical plate formation, respectively. Later-appearing Reelin-positive interneurons (Alcántara et al., 1998; Pesold et al., 1999; Meyer et al., 2002) may also be able to compensate for the loss of CR cells.

The intense Reelin signal in CR cells is thus redundant for lamination and architecture of the mouse cortex. However, to judge from the prominence of CR cells in the human brain (Meyer and Goffinet, 1998; Meyer et al., 1999, 2002), they gain in importance during evolution and may be involved in other aspects of cortical development, such as size increase and folding of the cortical surface. These aspects are not manifest in the lissencephalic mouse cortex but may be relevant for the formation of the gyrated primate cortex. The loss of the hippocampal fissure in the  $p73^{-/-}$  mice supports a possible relationship between p73 expression in CR cells and cortical folding.

Although p73-deficient mice do not have *reeler*-like malformations, cortical development is abnormal, and the ectopic cells in the preplate reflect an early alteration of the developmental program. The cortical hypoplasia points to a generalized defect of cell proliferation in the ventricular zone attributable to the loss of p73 and indicates that the activities of the p73 gene in brain development are wider than expected from the restricted expression in CR cells.

## The $p73^{-/-}$ hippocampus

The embryonic  $p73^{-/-}$  hippocampus has three distinctive features: absence of CR cells, lack of the hippocampal fissure, and dysgenesis of the dentate gyrus. Hippocampal CR cells and Reelin are essential for the establishment of the entorhinal-hippocampal pathways (Del Rio et al., 1997; Super et al., 1998; Ceranik et al., 2000), and the connectivity of the  $p73^{-/-}$  hippocampus is probably profoundly disturbed. We show that CR cells may also be critical for the folding of the hippocampal fissure. Because the fissure develops normally in reeler mice, folding does not depend on Reelin; however, CR cells may play a role in this process via p73 or other proteins, e.g., the chemokine receptor CXCR4 (Stumm et al., 2003; Tissir et al., 2004), whose ligand, SDF1 (stromal cell-derived factor 1), is highly expressed in the meninges and in cells investing the hippocampal fissure (Lu et al., 2002). Meningeal cells strongly influence dentate development, and their destruction induces a loss of the infrapyramidal blade (Hartmann et al., 1992) similar to the  $p73^{-/-}$  dentate defect.

On the other hand, it has been proposed that subpial astrocytes in the fissure form a transient boundary between Ammon's horn and the dentate gyrus that allows the establishment of an independent radial glia scaffold in the dentate subgranular zone (Sievers et al., 1992). The radial glia scaffold is essential for the migration of dentate granule cells and defective in reeler and scrambler mice (Forster et al., 2002; Frotscher et al., 2003; Weiss et al., 2003). We show that it is also reduced and disorganized in  $p73^{-/-}$  mice. Importantly, astrocytes in the subgranular zone are also progenitors of late-born granule cells (Seri et al., 2001). Loss or reduction of this precursor population may lead to a decrease in adult neurogenesis, which has been correlated with learning and memory (Kempermann et al., 1997; Gould et al., 1999). In summary, CR cells, meninges, subpial astrocytes, and dentate radial glia are closely interconnected in hippocampal morphogenesis, and p73 is deeply involved in this process.

# The $p73^{+/-}$ brain phenotype

The  $p73^{+/-}$  malformation is mainly a patterning defect of the posterior pole. Areal territories are altered, with ventral regions (posterior cortical nucleus of the amygdala and entorhinal cortex) dominating over dorsal ones (posterior temporal and occipital neocortex). The neocortex is reduced and transformed into an entorhinal-like cortex. Furthermore, cell numbers in the caudal  $p73^{+/-}$  cortex continue to decrease postnatally by apoptosis (Pozniak et al., 2002). These abnormalities indicate a decisive role of p73 in the areal patterning and neuronal survival of the posterior telencephalon.

p73 heterozygosity affects number and position of the Reelin-positive stellate cells of entorhinal layer II, which are a major origin of the entorhinal-hippocampal pathway (Steward and Scoville, 1976; Witter, 1993). The ectopic Reelin-positive cells in the altered  $p73^{+/-}$  neocortex possibly establish aberrant connections. Another important aspect of the  $p73^{+/-}$  brain is the size reduction and transformation of visual and secondary auditory areas, which raises the question of whether the sensory pathways are correctly established. Remarkably,  $p73^{+/-}$  mice display apparently normal behavior despite their cortical defect. However, it is questionable whether a  $p73^{+/-}$ -like phenotype would be compatible with normal cognitive functions in the human brain. Extrapolation of the mouse brain defect on the human brain would predict a substantial reduction of primary visual cortex and occipitotemporal association areas, a damage that, combined with the entorhinal malformation, would lead to severe cognitive disabilities and mental retardation.

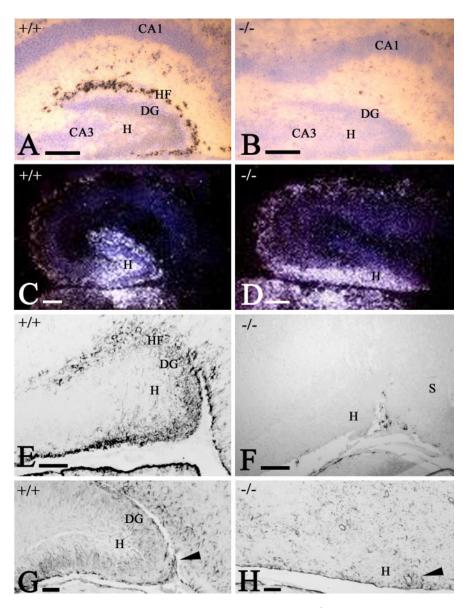
A comparable dosage-dependent defect has been described for the neurodevelopmental control gene PAX6, homozygous mutation of which is lethal in mice and humans. PAX6 heterozygous individuals present aniridia and behavioral and cognitive defects associated with structural abnormalities of specific cortical areas (Sisodiya et al., 2001; Ellison-Wright et al., 2004).

# An E2F1–TAp73 signaling pathway in the cortical hem

We reported previously that CR cells derive from the cortical hem and migrate tangentially into the neocortex (Meyer et al., 2002; Takiguchi-Hayashi et al., 2004). The cortical hem is a posteromedial signaling center that releases signaling molecules such as BMP (bone morphogenetic) and WNT (winglessint) proteins, which control the expression of region-specific tran-

scription factors; disruption of the signaling pathways in the hem results in distorted cortical and hippocampal patterning (Ragsdale and Grove, 2001; Grove and Fukuchi-Shimogori, 2003). Our proposal that early alterations of p73 expression in the hem lead to alterations of the posterior cortical map is consistent with this concept, although the possible molecular mechanisms are as yet unknown.

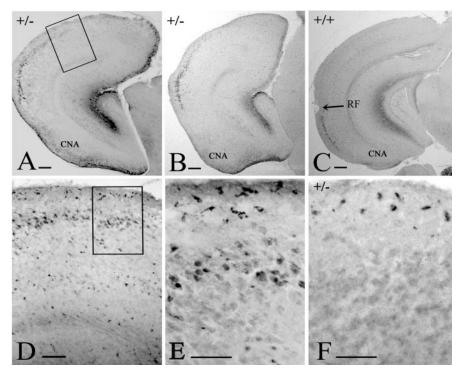
Does the role of p73 in cortical patterning depend on CR cells? Our findings point to a CR cell-independent mechanism. The severe malformation of the posterior pole in the  $p73^{-/-}$  mice cannot be attributed to CR cells because they are lost throughout the cortex, not only in the most affected region. On the other hand, CR cells are apparently unchanged in  $p73^{+/-}$  and  $E2F1^{-/-}$  mice, both of which present abnormal caudal regionalization. The differential expression of the main p73 products may give a



**Figure 7.** The postnatal hippocampal fissure (HF). A, B, Reelin ISH in WT (A) and  $p73^{-/-}$  (B) mice at P2. B, CR cells are absent, and the molecular layers of dentate gyrus (DG) and CA1/CA2 are fused. C, D, Calretinin ISH in WT (C) and  $p73^{-/-}$  (D) mice at P2. In the mutant, calretinin-positive hilus (H) cells have abnormal distribution. E, E, GFAP IHC in WT (E) and E0 and E1 mice at P2. GFAP stains subpial astrocytes in the HF and the incipient radial glia scaffold in the dentate gyrus. In the mutant, astrocytes are found only along the medial brain surface. E1, E2 for E3 in E4 mice at P12. The radial glia scaffold is fully established in the WT dentate; in the mutant, astrocytes are sparse and lack radial orientation. Arrowheads mark the entrance of the fissure, which fails to fold in the mutant. E3, Subiculum. Scale bars: E4, E7, E8, 200 E7, E8.

clue for solving this question because CR cells express DNp73 (Yang and McKeon, 2000; Yang et al., 2000), whereas TAp73 transcripts are detected in the cortical hem.

The *E2F1*<sup>-/-</sup> mice may help to shed light on the p73 patterning defect. E2F1 is a member of a transcription factor family that functions as a positive regulator of apoptosis and a suppressor of cell-cycle progression (Helin et al., 1992; Kaelin et al., 1992; Shan et al., 1992; Field et al., 1996). E2F1-deficient mice appear to develop normally, although postnatal and adult neurogenesis is impaired in olfactory bulb, dentate gyrus, and cerebellum (Cooper-Kuhn et al., 2002). E2F1 is a major activator of the TAp73 promoter (Irwin et al., 2000; Seelan et al., 2002), which is consistent with the significant decrease of TAp73 protein in the *E2F1*<sup>-/-</sup> brain. In contrast, DNp73 seems to be less affected by E2F1 deficiency than TAp73, which correlates with the preserved



**Figure 8.** Architecture of the  $p73^{+/-}$  telencephalon. Reelin in P3  $p73^{+/-}$  (A), P12  $p73^{+/-}$  (B), and P12 WT (C) brains. Reelin-IR cells in layer II mark the entorhinal cortex in the WT brain. In  $p73^{+/-}$  brains, they are dorsally displaced and expanded, whereas the neocortex is reduced in size. The posterior cortical nucleus of the amygdala (CNA) is hyperplastic. The rhinal fissure (RF) is not visible in the  $p73^{+/-}$  brains. D, E, Higher magnification of the boxed areas in E and E0, respectively, to show ectopic Reelin-IR cells in layer II of the abnormal cortex and CR cells in layer I. E7, At P3, CR cells in the abnormal E7, cortex are intensely p73 IR. Scale bars: E7, 200 E8, E9, 100 E9, 1

p73 immunoreactivity in CR cells. E2F1 has other targets in addition to TAp73 (Stanelle et al., 2002), and, in fact, the  $E2F1^{-/-}$  phenotype is not completely identical to the  $p73^{+/-}$  phenotype. The rather mild defect in  $E2F1^{-/-}$  mice and the more severe form in the  $p73^{+/-}$  mice point to a dosage-dependent effect of TAp73 in cortical patterning.

Our data in the p73 and E2F1 mice suggest that the malformation of the posterior pole is unrelated to CR cells. The common feature in  $p73^{-/-}$ ,  $p73^{+/-}$ , and  $E2F1^{-/-}$  brains is the loss and reduction, respectively, of TAp73 expression, supporting the hypothesis that an E2F1–TAp73 pathway in the cortical hem is important for the regionalization of the posterior pole.

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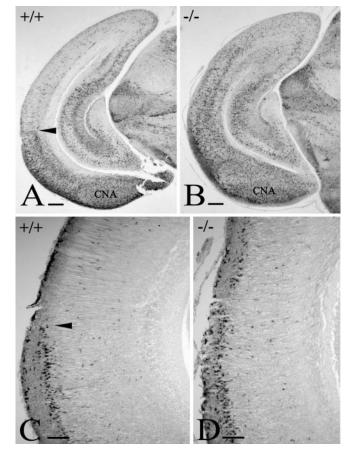
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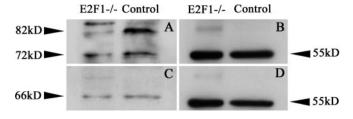
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**Figure 9.** The  $E2F1^{-/-}$  cortex. Calretinin immunostaining in WT (A) and  $E2F1^{-/-}$  (B) cortex at P1 shows that the caudal neocortex is transformed and acquires entorhinal-like features. Calretinin-IR interneurons are distributed differently in WT and  $E2F1^{-/-}$  cortex, and the posterior cortical nucleus of the amygdala (CNA) is slightly larger in the mutant. C, In WT, Reelin-IR cells in layer II stop at the rhinal fissure (arrowheads in C and C). C, in the mutant, they extend dorsally into the altered neocortex. Scale bars: C, C, C, C, C0, C0 C0.



**Figure 10.** Expression of p73 in E2F1 knock-out mice. A, C, Immunoblot analysis of p73 in cortical tissue extracts probed with N-terminal (A) and C-terminal (C) antibodies. Note the important reduction of TAp73 levels at 82 kDa (TAp73 $\alpha$ ) and at 72 kDa (probably representing TAp73 $\beta$ ) in mutant mice compared with the controls. After incubation with a C-terminal antibody, a DNp73 isoform at 66 kDa shows only a slight reduction in the E2F1 mutant (C). C, C, C, C. Tubulin controls to test the amount of protein analyzed.

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