



# PAF-AH catalytic subunits modulate the Wnt pathway in developing GABAergic neurons

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Platelet-activating factor acetylhydrolase 1B (PAF-AH) inactivates the potent phospholipid platelet-activating factor (PAF) and is composed of two catalytic subunits ( $\alpha 1$  and  $\alpha 2$ ) and a dimeric regulatory subunit, LIS1. The function of the catalytic subunits in brain development remains unknown. Here we examined their effects on proliferation in the ganglionic eminences and tangential migration. In  $\alpha 1$  and  $\alpha 2$  catalytic subunits knockout mice we noticed an increase in the size of the ganglionic eminences resulting from increased proliferation of GABAergic neurons. Our results indicate that the catalytic subunits act as negative regulators of the Wnt signaling pathway. Overexpression of each of the PAF-AH catalytic subunits reduced the amount of nuclear beta-catenin and provoked a shift of this protein from the nucleus to the cytoplasm. In the double mutant mice, Wnt signaling increased in the ganglionic eminences and in the dorsal part of the cerebral cortex. *In situ* hybridization revealed increased and expanded expression of a downstream target of the Wnt pathway (*Cyclin D1*), and of upstream Wnt components (*Tcf4*, *Tcf3* and *Wnt7B*). Furthermore, the interneurons in the cerebral cortex were more numerous and in a more advanced position. Transplantation assays revealed a non-cell autonomous component to this phenotype, which may be explained in part by increased and expanded expression of *Sdf1* and *Netrin-1*. Our findings strongly suggest that PAF-AH catalytic subunits modulate the Wnt pathway in restricted areas of the developing cerebral cortex. We hypothesize that modulation of the Wnt pathway is the evolutionary conserved activity of the PAF-AH catalytic subunits.

**Keywords:** platelet-activating factor acetylhydrolase 1B, Wnt, beta-catenin, ganglionic eminences

## INTRODUCTION

The proper functioning of the cerebral cortex relies on formation of neural networks that are composed of excitatory neurons and inhibitory interneurons (reviewed by Wonders and Anderson 2006). The excitatory, or projection, neurons are born in proliferating zones of the cerebral cortex, the ventricular zone and the subventricular zone (Noctor et al., 2004; review Gotz and Huttner, 2005). These neurons usually migrate along radial glia to their proper cortical layer (reviews by Hatten, 2002; Ayala et al., 2007). The majority of interneurons, which compose approximately 20% of cortical neurons, are born in the ganglionic eminences (GE) and migrate to the cortex using a tangential mode of migration (reviews by Marin and Rubenstein, 2001; Metin et al., 2006). Following their arrival to the cerebral cortex, the interneurons utilize radial migration to reach the proper laminar position and then they intercalate in the network. Gamma-aminobutyric acid (GABA)-biosynthesizing enzymes preferentially localize to cortical interneurons, which are also known as GABAergic neurons. In the adult mammalian brain, GABA has been associated primarily with the mediation of synaptic inhibition. Imbalance between inhibition and excitation may underlie diseases such as epilepsy (reviewed by Ben-Ari, 2006). Cortical interneurons have also been implicated in developmental processes, including the regulation of neuronal proliferation and migration during corticogenesis and the development of cortical circuitry (reviews by Owens and Kriegstein,

2002; Spitzer, 2006). Identification of molecules and pathways, which regulate the number and migration of GABAergic neurons, is of clear importance when aiming to understand the normal and diseased brain.

Platelet-activating factor-acetyl hydrolase (PAF-AH) hydrolyzes PAF, an important lipid second messenger. PAF mediates an array of biological processes and is very abundant in the mammalian central nervous system, where it acts as a synaptic messenger, a transcription inducer and is involved in long-term potentiation, a cellular model of memory formation (Kato et al., 1994; Bazan, 1995). PAF deacetylation by PAF-AH leads to its inactivation. The PAF-AH1B intracellular enzyme is a tetramer composed of two catalytic subunits,  $\alpha 1$  (PAFAH1B3) and  $\alpha 2$  (PAFAH1B2), and a regulatory dimer of LIS1 (PAFAH1B1) (Hanahan, 1986; Kornecki and Ehrlich, 1988; Koltai et al., 1991; Chao and Olson, 1993; Stafforini et al., 2003; Bazan, 2005). From here forth PAF-AH1B will be referred to as PAF-AH. Deletions or point mutations in the *LIS1* (*Lissencephaly-1*) gene are manifested in humans in a spectrum of abnormal neuronal migration phenotypes ranging between classical lissencephaly and subcortical band heterotopia in humans (Reiner et al., 1993; Barkovich et al., 2005). Impaired neuronal migration has been noticed also in mouse genetic models (Hirotsume et al., 1998; Cahana et al., 2001; McManus et al., 2004b). Further reduction of LIS1 levels using *in utero* electroporation or conditional knock-out results in further impairment of neuronal

migration as well as affects the proliferation of progenitors (Shu et al., 2004; Tsai et al., 2005; Tsai et al., 2007; Hebbar et al., 2008; Yingling et al., 2008). LIS1 dosage is crucial for proper brain development an increased levels of LIS1 affect brain development in human and in mice (Bi et al., 2009). Increased LIS1 levels affect the organization of the ventricular zone and impaired radial and tangential neuronal migration.

Platelet-activating factor-acetyl hydrolase catalytic subunits knockout mice were generated independently by two groups (Koizumi et al., 2003; Yan et al., 2003). Deletions of only the  $\alpha 2$  subunit or both the  $\alpha 1/\alpha 2$  subunits resulted in severe male infertility. No brain phenotype has been reported for these knockout mice. We further investigated these mice (Koizumi et al., 2003) and detected a moderate increase in the size of the ganglionic eminences. In parallel, we noted that overexpression of the PAF-AH catalytic subunits modulated the Wnt pathway in transfected cells. Conversely, in the double knockout mice Wnt signaling was enhanced. An additional effect on tangential migration was noted: the interneurons in the cerebral cortex were more numerous and in a more advanced position, which was accompanied with earlier development of thalamocortical fibers. Our findings strongly suggest that PAF-AH catalytic subunits modulate the Wnt pathway in restricted areas of the developing cerebral cortex.

## MATERIALS AND METHODS

### AUTHORIZATION FOR THE USE OF EXPERIMENTAL ANIMALS

The experiments described in this manuscript were approved by the Weizmann Institute IACUC.

### PLASMIDS

The expression construct of pDsRed $\alpha 1$  was generated using rat  $\alpha 1$  tagged with green fluorescent protein (GFP) digested with *EcoR1* and *NotI*. The insert was cloned between *Xho1* and *SmaI* with the *EcoR1*–*Xho1* adaptors (5′-TCGAGCTACCTGCGGTG-3′ and 5′-AATTCACCGCAGGTAGC-3′) into pDsRed-C1 vector (Clontech, CA, USA). The corresponding Rat  $\alpha 2$  tagged with GFP was digested with *EcoR1* and *Kpn1* and cloned between *Xho1* and *Kpn1* using the same adaptors and vector as described above. pcDNA3-Flag $\alpha 1$  and pcDNA3-Flag $\alpha 2$  expression vectors were generated by subcloning the same fragments into pcDNA3 tagged with a Flag epitope. Cloning was verified by DNA sequence analysis of the plasmids, and Western blot analysis of the transfected expression plasmids in mammalian cells. A plasmid that contains the 3′ UTR of the  $\alpha 1$  was generated using a clone isolated from mouse brain cDNA library, inserted into pBS following *PstI* digestion. pcDNA3-Myc $\alpha 1$  plasmid was received from Prof. Junken Aoki. All GFP- $\beta$ -catenin constructs, pTOPFLASH-Luc and pFOPFLASH-Luc were kindly provided by Prof. Avri Ben-Ze'ev. pCIG-CA- $\beta$ -catenin plasmid was a gift from Prof. Andrew P. McMahon (Megason and McMahon, 2002). Flag-Dvl construct was a kind gift from Prof. Marianne Bienz (Schwarz-Romond et al., 2005). pTLHA-CRMP1 was provided by Prof. Erich E. Wanker (Stelzl et al., 2005) and pCDNA-HA-TLE1 was received from Prof. Yoram Groner (Levanon et al., 1998). CXCL12 (*Sdf-1*), inserted into pCMV-SPORT6 vector, was a gift from Prof. John L.R. Rubenstein. pBSc- $\alpha 2$  and *Cyclin D1*, *Tcf3*, *Tcf4* and *Wnt7B* in pFLCI vectors, were provided by the Forchheimer plasmid collection.

### ANTIBODIES

Monoclonal antibodies specific for Myc (SC-40) and HA (16B12) were purchased from Santa Cruz, CA, USA, and Constance CA, USA, respectively. Mouse monoclonal anti-Flag (M2), polyclonal antibodies specific for  $\beta$ -catenin and Goat anti-mouse-tubulin (DM1A) were purchased from Sigma (Rehovot, Israel). Mouse monoclonal antibodies for pY489- $\beta$ -catenin were obtained from the developmental studies hybridoma bank (DSHB). Rabbit polyclonal antibodies for Calbindin were purchased from SWANT (Bellinzona, Switzerland). Polyclonal antibodies specific for Cyclin D1 were purchased from Abcam Cambridge, UK. TAG-1 antibodies were received from Prof. Peles (Horresh et al., 2008). Secondary antibodies used for immunostaining included Cy3- and Cy5-conjugated donkey anti-mouse and anti-rabbit IgG (H + L), and Cy3-conjugated Goat anti-mouse IgM purchased from Jackson ImmunoResearch.

### CELL LINES

COS-7 and HEK293 were grown at 37°C, 5% CO<sub>2</sub> in DMEM (Gibco, Auckland, New Zealand) supplemented with 10% fetal bovine serum (FBS), 4 mM glutamine, 100 U/ml penicillin and 0.1 mg/ml streptomycin.

### MICE

The single KO PAF-AH catalytic subunit mice were received from Prof. Hiroyuki Arai (Koizumi et al., 2003) and were bred to generate the double KO mice. Prof. Yuchio Yanagawa kindly provided the GAD67-GFP ( $\Delta$ neo) mice (Tamamaki et al., 2003). Mice were raised in the Weizmann Institute of Science transgenic facility. Staging of embryos were according to defined developmental stages (Kaufman, 1992). The GAD67-GFP mice were bred with the double KO mice to create triple transgenic mice.

### TRANSFECTIONS

COS-7 Cells were transfected using JET PEI (Polyplus-transfection, NY, USA). HEK293 cells were transfected by the calcium phosphate precipitation method (Graham and van der Eb, 1973).

### LUCIFERASE REPORTER ASSAYS

Lef/Tcf reporter assays were performed as previously described (Brembeck et al., 2004). Briefly, HEK293 cells were trypsinized 1 day prior to transfection and plated on 24 well plates. Cells were co-transfected with 0.25  $\mu$ g of pTOPFLASH-Luc or pFOPFLASH-Luc, either 0.5  $\mu$ g of pCIG-CA- $\beta$ -catenin or Dvl expression constructs, and either TLE1 or CRMP1, respectively and also with different combinations of either PAF-AH  $\alpha 1$  or  $\alpha 2$ . Empty pcDNA3 was added to keep the plasmid amounts equal in each transfection. Luciferase activity was determined 48 hr after transfection and normalized against  $\beta$ -galactosidase activity. Reporter assays were performed as triple transfections.

### IMMUNOSTAINING

COS-7 cells were grown on coverslips (Menzel-Glaser, Braunschweig, Germany). Two days after transfection, cells were washed in PBS (Gibco, Auckland, New Zealand), fixed with 3% paraformaldehyde and permeabilized using 0.1% Triton X-100. After blocking three times in PBS supplemented with 0.1% BSA (Sigma, Rehovot,

Israel), coverslips were incubated with the indicated antibodies, stained with DAPI, mounted with Immu-mount (Thermo Electron Corporation, USA) and examined using the DeltaVision system (Applied Precision, Issaquah, WA, USA).

### BRAIN IMMUNOSTAININGS

E13.5 wild-type, heterozygote ( $\alpha 1^{+/-}\alpha 2^{+/-}$ ), or double KO ( $\alpha 1^{-/-}\alpha 2^{-/-}$ ) embryos were fixed in 4% paraformaldehyde overnight. The post-fixed embryos were washed three times with PBS and brains were dissected out. The brains were then embedded in 3.5% low-temperature melting point agarose. Coronal sections (60  $\mu$ m) were cut using a vibratome (Leica) and collected floating in PBS. Sections were permeabilized with PBS-T (1  $\times$  PBS, 0.1% Triton X-100), blocked in 10% FCS/PBS-T and then incubated overnight at 4°C with the indicated antibodies. Slides were washed with PBS-T and incubated with secondary antibodies for 1 h RT, washed with PBS-T and mounted using Aqua Poly/Mount (Polysciences, Inc., USA).

### IN SITU HYBRIDIZATION

E13.5 embryos (wild-type C57BL/6J, Heterozygote for  $\alpha 1$  and  $\alpha 2$  ( $\alpha 1^{+/-}\alpha 2^{+/-}$ ), or double KO ( $\alpha 1^{-/-}\alpha 2^{-/-}$ ) were collected in ice-cold PBS in a RNase-free environment, then fixed in 4% paraformaldehyde overnight. Isolated brains were gradually dehydrated to 100% methanol and stored at -20°C. After bleaching with 1:5 H<sub>2</sub>O<sub>2</sub>:methanol, the brains were rehydrated and treated with Proteinase K, then brains were re-fixed and sectioned using a Leica vibratome (150  $\mu$ m). Antisense digoxigenin-labeled RNA templates were generated by *in vitro* transcription using PCR products (using the appropriate combinations of T7, T3, and SP6 primers) from the corresponding genes. Hybridization was conducted overnight at 60°C, using the riboprobes at a concentration of 1 ng/ $\mu$ l in hybridization buffer [50% formamide (Ambion, Israel), 1.3% SSC, 0.2% Tween-20, 5 mM EDTA, 50  $\mu$ g/ml yeast tRNA, 100  $\mu$ g/ml Heparin (Sigma, Israel)]. Immunological detection was done using anti-digoxigenin-AP-conjugate antibodies (Roche, Germany) and NBT/BCIP reagents (Boehringer, Germany).

### TRANSPLANTATION TO EXPLANTS

E13.5 GAD67-GFP, wild-type and double KO brains were cut into 250  $\mu$ m coronal slices and then transferred onto inserts (MilliCell-CM, Millipore) floating on serum free medium (Neurobasal medium supplemented with B27, N2, GlutaMax, glucose, and gentamicin). MGE explants, dissected from E13.5 GAD67-GFP brains, were loaded into a glass micropipette and transplanted into either wild-type or double KO E13.5 MGE. After 24 h of incubation at 37°C, 5% CO<sub>2</sub>, grafted neurons were visualized using the DeltaVision system package. Each time-lapse movie lasted 3 h with one frame taken every 2 min.

### DiI LABELING

To examine thalamocortical projections, fixed brains (E13.5) were labeled with DiI crystals (1,1'-dioctadecyl-3,3',3'-tetramethylindocarbocyanine perchlorate ('DiI'; DiI18(3)) \*crystalline\*, Molecular probes). The crystals were placed within the thalamus and were left to diffuse for 5 days prior to vibratome sectioning (60  $\mu$ m) that were later imaged using standard fluorescent microscopy.

### MICROSCOPY

Images were taken using either the DeltaVision system package (Applied Precision, Issaquah, WA, USA), or confocal microscopy (LSM510, Zeiss).

### MEASUREMENTS

Cell counts and neuronal migration speed were analyzed using the MeasurementPro and ImarisTrack modules of Imaris software (Bitplane, Zurich, Switzerland). Area was measured using Adobe Photoshop CS2.

### STATISTICAL ANALYSIS

Statistical analysis was conducted using Prism 4 for Macintosh (GraphPad Software, Inc.).

## RESULTS

### DOUBLE MUTANT MICE EXHIBIT INCREASED GE AREA, WHERE PAF-AH CATALYTIC SUBUNITS ARE EXPRESSED

Analysis of the brains of double mutant embryos (E13.5), revealed a modest increase in the size of the ganglionic eminences (GE). The area of the GE and number of cells at M-phase were determined by anti-phosphohistone H3 (pH3) immunostaining (Figures 1A,B). Double KO mice had a significant ( $P = 0.0068$ )  $34 \pm 9.4\%$  increase ( $112600 \pm 7579$  pixels,  $n = 8$ ) in the area of pH3 positive cells in comparison with the heterozygotes ( $83860 \pm 2371$  pixels,  $n = 10$ ). In addition, double KO mice exhibited a  $26 \pm 7.4\%$  increase in the number of pH3 positive cells ( $336.4 \pm 18.9$ ,  $n = 8$ ,  $P = 0.0029$ ) in comparison with the heterozygotes ( $266.9 \pm 9.3$ ,  $n = 10$ ). When we examined the number of pH3 positive cells in the cerebral cortex there was no difference between the double KO and the heterozygotes, thus suggesting an area specific effect of the mutation. When E12.5 embryos were examined, the area of the proliferation zone in the GE was only slightly and not significantly increased (+4.8%). There was no difference in the number of pH3 positive cells, therefore we focused our studies on E13.5 embryos.

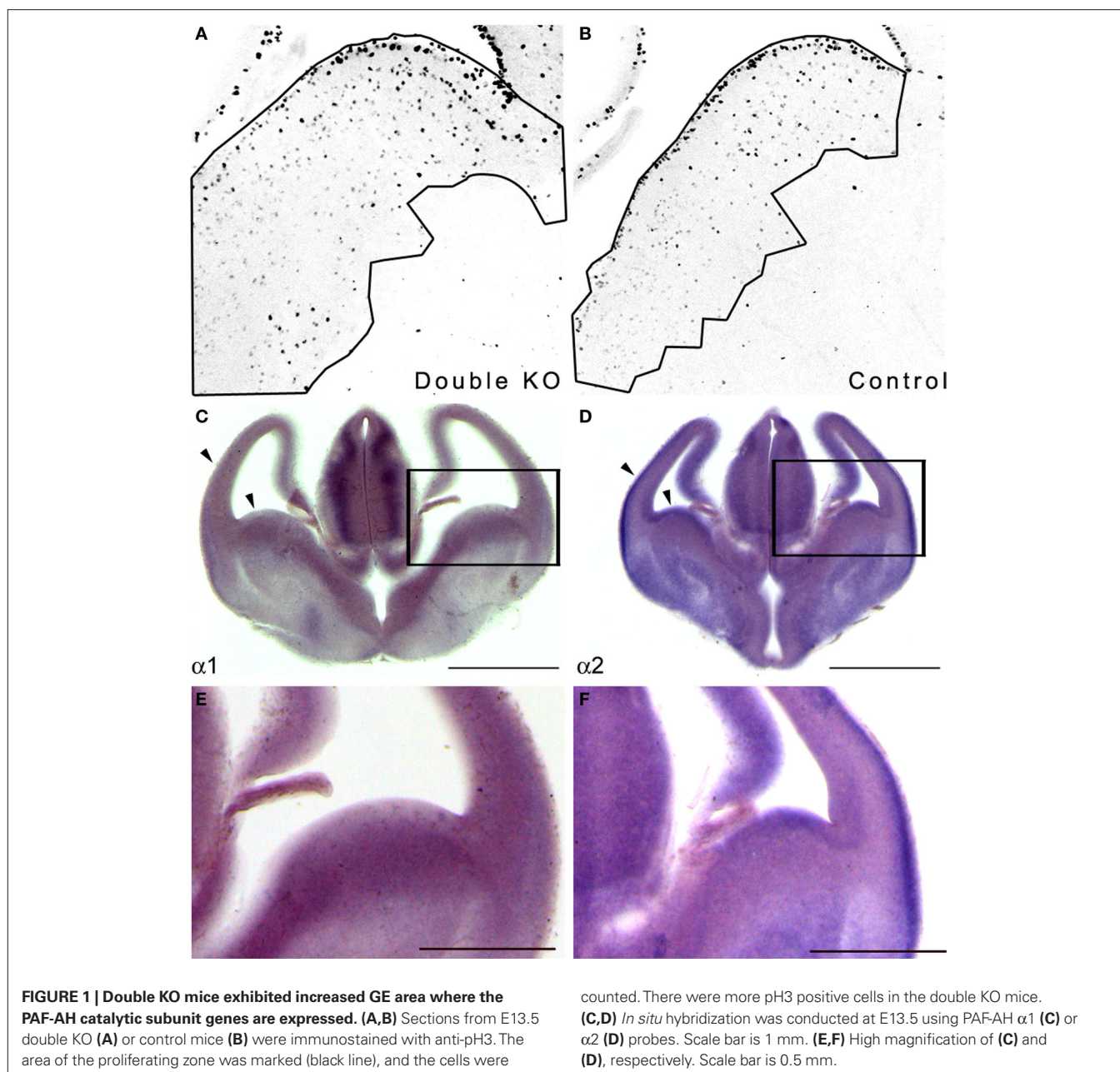
Next, we examined whether the PAF-AH subunits are expressed at this developmental stage in the mouse embryonic brain. *In situ* hybridization showed that both the  $\alpha 1$  and the  $\alpha 2$  PAF-AH subunits are expressed in the GE as well as in other brain regions such as the developing cerebral cortex and the thalamus (Figures 1C-F).

Embryonic expression of the PAF-AH catalytic subunits has been previously noted in the mouse and rat brain (Albrecht et al., 1996; Many et al., 1998). Our results suggest that the increased number of proliferating cells in the GE observed in double KO at E13.5 may be associated with the lack of expression of PAF-AH catalytic subunits.

### PAF-AH SUBUNITS MODULATE THE Wnt PATHWAY

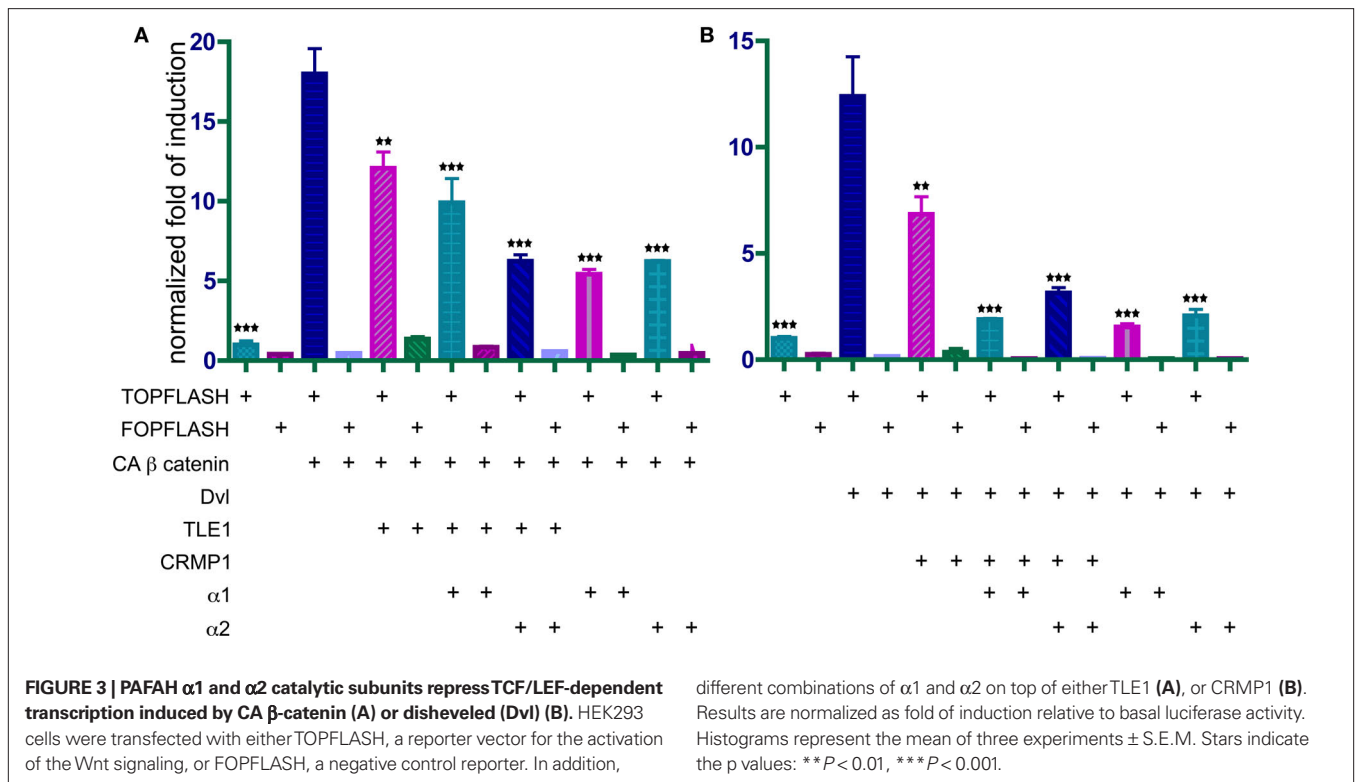
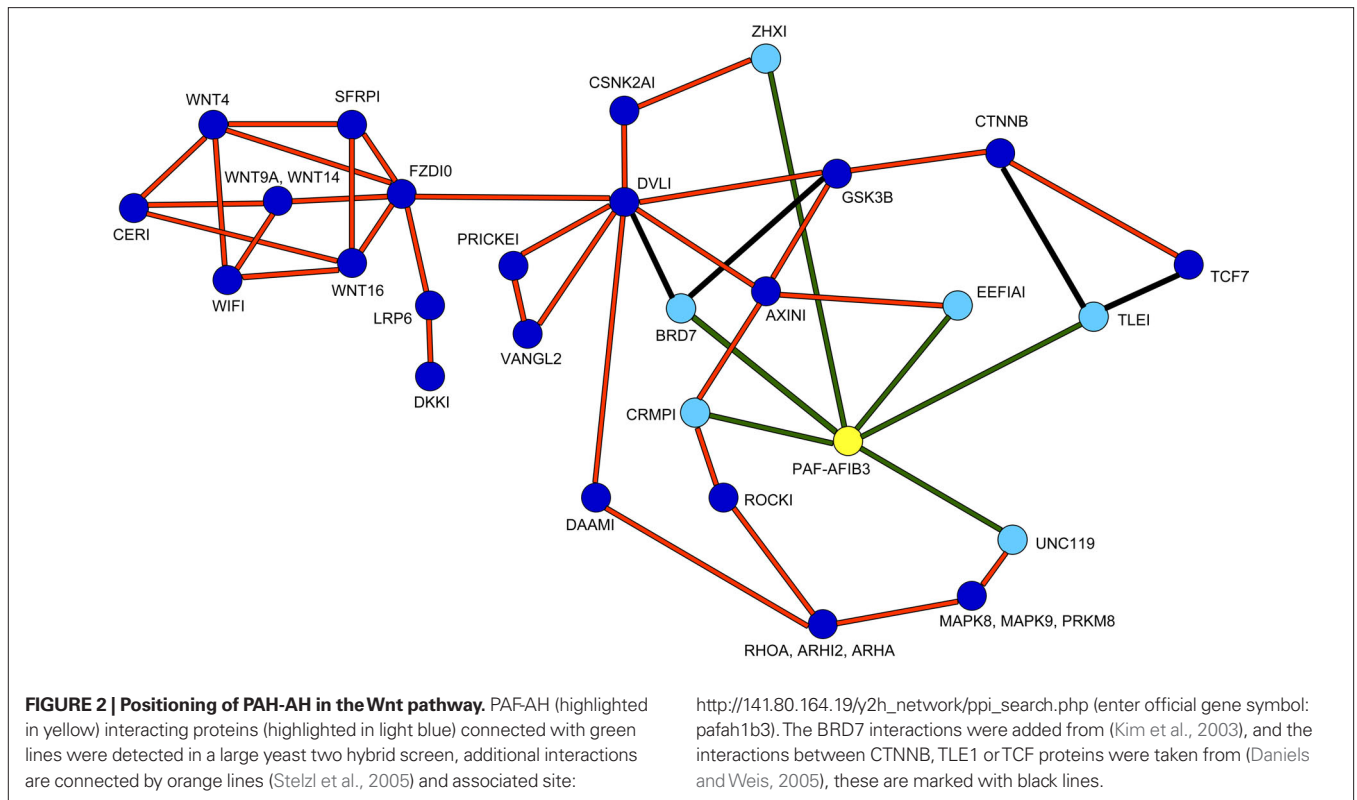
The Wnt pathway participates in the regulation of neuronal proliferation in the ganglionic eminences (Gulacsi and Anderson, 2008). Furthermore, high throughput protein-interaction data suggested possible interactions of the PAF-AH 1B3 ( $\alpha 1$ ) catalytic subunit with several proteins related to the Wnt pathway (Kim et al., 2003; Daniels and Weis, 2005; Stelzl et al., 2005). A schematic presentation of the suggested position of PAF-AH within the Wnt pathway is shown in Figure 2.





As can be seen in the scheme, PAF-AH interacts with multiple components both up and downstream to the canonical Wnt pathway. Although we verified some of the physical interactions in transfected cells using co-immunoprecipitation assays (CRMP1, BRD7 and UNC119, data not shown), we still could not predict how modulation of PAF-AH subunits will affect Wnt-mediated signaling in regard to modulating upstream and/or downstream processes. To test the possible effect of the PAF-AH subunits on the Wnt pathway we conducted cell-based luciferase reporter assays (Figure 3). Activation of the Wnt pathway stabilizes  $\beta$ -catenin allowing its entry to the nucleus, thereby activates transcription of genes containing TCF/LEF1 binding sites (Eastman and Grosschedl, 1999). The activation can be induced by addition of a constitutively active (CA) form of  $\beta$ -catenin

or by addition of a Disheveled (Dvl) protein. HEK293 cells were co-transfected with either TOPFLASH (positive reporter plasmid with three copies of TCF/LEF-1 binding sites) or FOPFLASH (negative control with mutated binding sites) reporters, along with either CA  $\beta$ -catenin or Dvl (Figures 3A,B, respectively). Overexpression of CA  $\beta$ -catenin or Dvl resulted in a significant increase in reporter gene activity ( $P < 0.001$ ). However, when either CRMP1 or TLE1 were co-expressed with Dvl or CA  $\beta$ -catenin, respectively, the increase was significantly smaller ( $P < 0.01$ ). A more pronounced reduction in the reporter activity was accomplished when either PAF-AH  $\alpha 1$  or  $\alpha 2$  proteins were expressed together with either CRMP1 or TLE1 ( $P < 0.001$ ). Nonetheless, it is important to note that a significant reduction in CA  $\beta$ -catenin- or Dvl-mediated activation was obtained



only when either  $\alpha 1$  or  $\alpha 2$  were co-expressed with these Wnt pathway activators ( $P < 0.001$ , ANOVA statistical analysis of luciferase assays with Tukey's multiple comparison test,  $n = 3$ ). Our results

verified that CRMP1 and TLE1 are negative regulators of the Wnt signaling pathway as previously reported (Levanon et al., 1998; Stelzl et al., 2005). More importantly, our results demonstrated that the

addition of either PAF-AH catalytic subunits  $\alpha 1$  or  $\alpha 2$  even without the addition of either CRMP1 or TLE1, repressed gene expression mediated by TCF/LEF1 binding sites.

In summary, expression of either PAF-AH catalytic subunit resulted in a decrease in activation of the Wnt-mediated reporter either following activation with CA  $\beta$ -catenin or following activation of the pathway using Dvl.

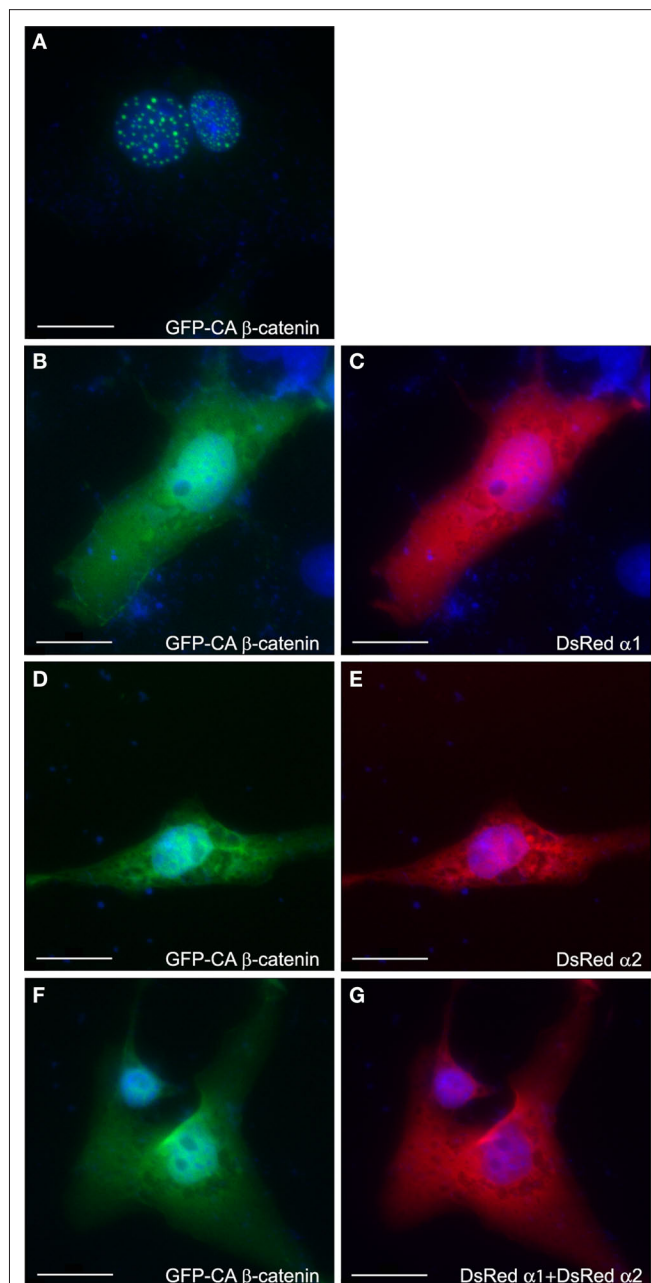
The intracellular localization of GFP-tagged CA  $\beta$ -catenin was used as an additional mean to monitor activation of the Wnt pathway. In transfected COS-7 cells CA  $\beta$ -catenin is typically found as puncta in the cell nucleus (**Figure 4A**  $92.3 \pm 6.2\%$  of the transfected cells) similar to previous reports (Simcha et al., 1998; Giannini et al., 2000). However, the intracellular localization of CA  $\beta$ -catenin shifted to a prominent cytoplasmic localization following transfection with either PAF-AH  $\alpha 1$  subunit, where only  $35.4 \pm 2\%$  of the transfected cells showed nuclear localization (**Figures 4B,C**) or PAF-AH  $\alpha 2$  subunit, where  $28.5 \pm 2.8\%$  of the transfected cells showed nuclear localization (**Figures 4D,E**). In the presence of both subunits a smaller proportion of cells exhibited nuclear localization of CA  $\beta$ -catenin (**Figures 4F,G**  $26.2 \pm 6.7\%$  of the cells exhibited nuclear localization). The reduction in the percentage of cells exhibiting nuclear  $\beta$ -catenin is statistically significant for all treatments (ANOVA analysis with Dunnett's multiple comparison test,  $P < 0.01$ ,  $n = 3$  experiments, 30 cells were quantified for each transfection in each experiment).

The apparent intracellular localization was also verified by cellular fractionation followed by Western blot analysis. Following addition of either PAF-AH  $\alpha 1$  or  $\alpha 2$  subunit there was a noticeable decrease in the proportion of nuclear  $\beta$ -catenin (data not shown). Interestingly, the catalytic subunits of PAF-AH were found in the nuclear as well as in the cytoplasmic fraction.

*In vivo* we noticed a dramatic expansion in the area expressing nuclear  $\beta$ -catenin, which was most pronounced in the mantle layer of the lateral cortex (**Figure 5**). There was an almost two fold increase in the number of cells exhibiting nuclear  $\beta$ -catenin staining in the double KO brains in comparison with control sections ( $192 \pm 34\%$  versus  $100 \pm 7.6\%$ ,  $n = 4$ ,  $P = 0.04$ ).

Next, we examined the expression pattern of additional components of the Wnt pathway using *in situ* hybridization (**Figure 6**). The expression pattern of *Cyclin D1*, a downstream target of the Wnt pathway, was increased and expanded in the double KO versus the control mice (**Figures 6A,B**).

*Tcf4* and *Tcf3* are members of the T cell factor/Lymphocyte enhancer factor family of transcription factors. Following activation of the Wnt pathway,  $\beta$ -catenin, alone or in association with members of the Tcf/Lef1, can translocate to the nucleus, where it ultimately controls the activity of specific target genes. We noticed an expansion in the expression domains of *Tcf4* (**Figures 6C,D**) and of *Tcf3* (**Figures 6E,F**). It is interesting to note that for the most, the expression pattern of *Tcf4* and *Tcf3* were mutually exclusive. In addition, a striking increase in *Wnt7B* expression was noticed mainly in the lateral cerebral cortex where we have previously noted an increase in nuclear  $\beta$ -catenin (**Figures 6G,H**, arrowhead). Collectively, our data indicate that the double KO mice exhibit increased and expanded expression of upstream and downstream Wnt-pathway components.

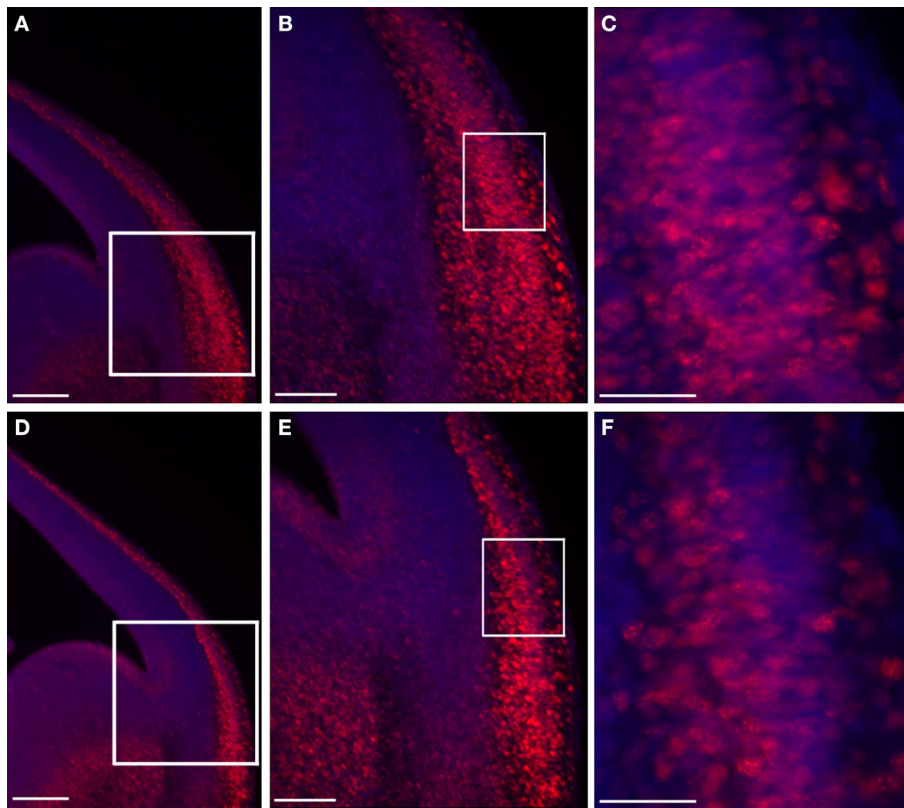


**FIGURE 4 | PAF-AH subunits affect subcellular localization of GFP-CA  $\beta$ -catenin.** COS-7 cells were transfected with GFP-CA  $\beta$ -catenin (**A**) alone or together with DsRed $\alpha 1$  (**B,C**) or DsRed $\alpha 2$  (**D,E**), or both PAF-AH catalytic subunits (**F,G**). Cells were fixed 48 h after transfection and nuclear DNA was stained with DAPI (shown in blue). Overexpression of GFP-CA  $\beta$ -catenin demonstrated nuclear localization (**A**), whereas the addition of either DsRed $\alpha 1$  (**C**) or DsRed $\alpha 2$  (**E**) or both (**G**) shifted  $\beta$ -catenin to the cytoplasm (**B,D,F**). Scale bars are 20  $\mu$ m.

#### PAF-AH DOUBLE KO MICE EXHIBIT ENHANCED TANGENTIAL MIGRATION

The development of the thalamocortical projection depends on the early tangential migration of a population of neurons derived from the ventral telencephalon (Lopez-Bendito et al., 2006). Therefore, we decided to examine whether the double KO





**FIGURE 5 | Double KO brain sections exhibit an increased area of nuclear  $\beta$ -catenin.** Double KO (A–C) exhibit more cells with nuclear  $\beta$ -catenin in comparison with control (B–F). The white boxes define the higher magnifications areas. Scale bars: (A,D) 200, (B,E) 100, (C,F) 40  $\mu$ m.

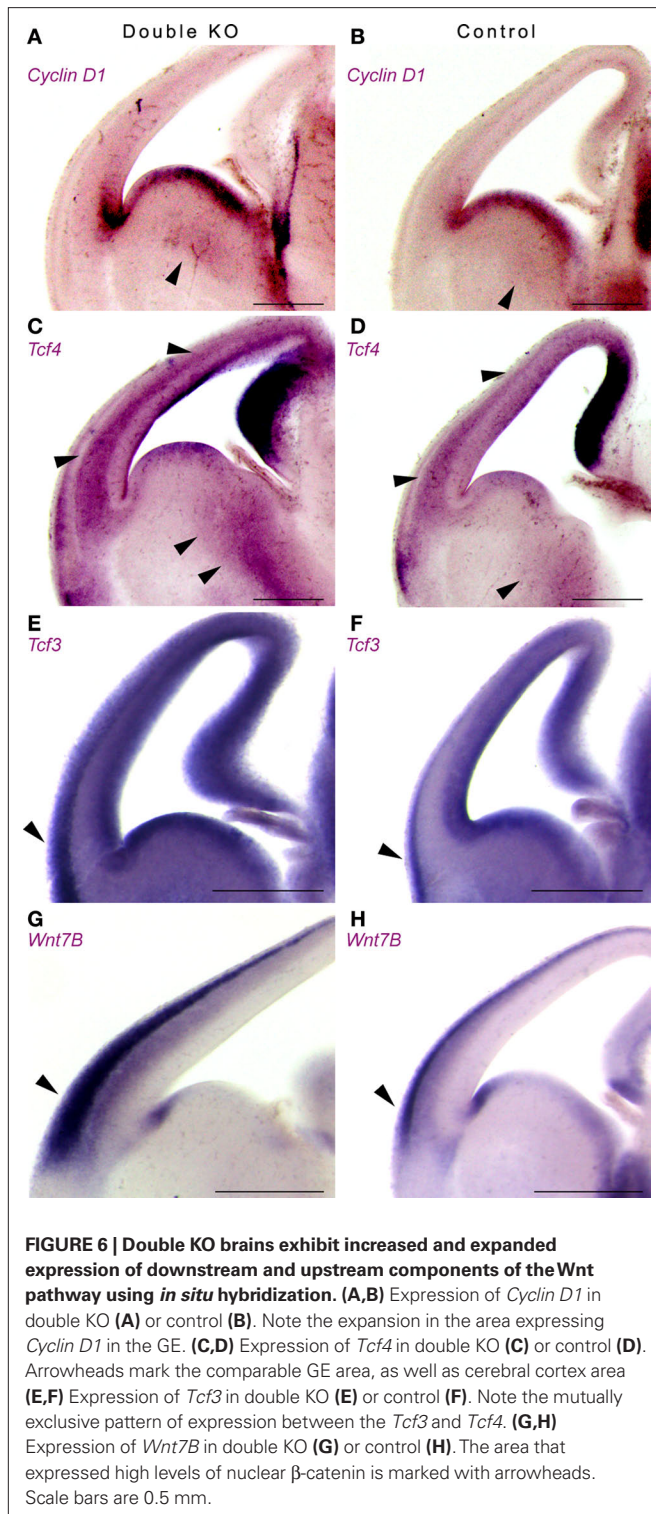
mice exhibit any changes in the development of thalamocortical projections. We used TAG-1 immunostaining to visualize the emergence of the corticofugal system in the developing cortex (Denaxa et al., 2001, 2005). Immunohistochemical staining of cortical sections of E13.5 mouse embryos revealed the presence of labeled cells and their axons in the region of the plexiform primordium in wild-type mice (Figures 7B,D). In the double mutant mice, intense TAG-1 immunoreactivity was present in fibers of the intermediate zone, the cortical plate as well as the marginal zone of the basolateral cortex (Figures 7A,C). In addition, the labeling of thalamocortical fibers by DiI revealed an early development of thalamocortical projections in the double KO brains (Figure 7E versus F).

The migration of interneurons from the ganglionic eminences towards the cortex was visualized by using fluorescent-based genetic labeling of these cells. We crossed GAD67-GFP knock-in mice (Tamamaki et al., 2003) with the double KO mice. An observable enhancement in interneuron migration to the cortex was noted following elimination of the PAF-AH catalytic subunits (Figures 8A,B versus C,D). These results were corroborated by an observable increase in the number and relative position of calbindin positive interneurons found in the cerebral cortex in the double KO in comparison with the control (Figures 8E,F versus G,H).

Next, we monitored neuronal migration by time-lapse movies of neurons. Explants of GAD67-GFP MGE neurons, transplanted on

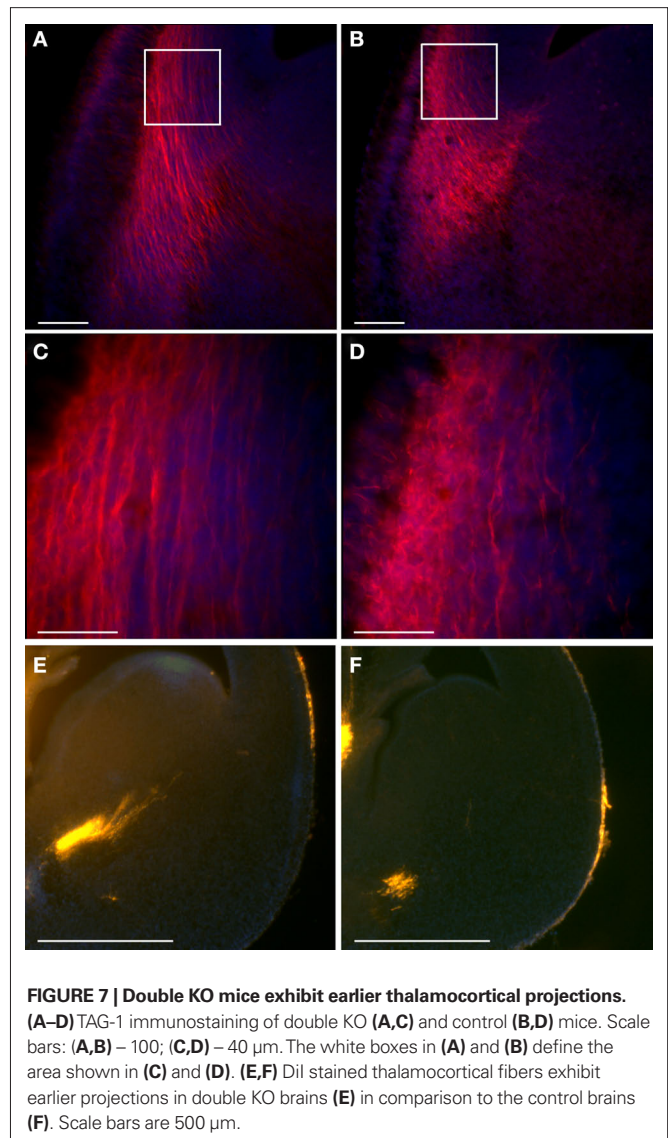
either a wild-type or double knockout substrate, were time-lapsed 24 h after transplantation. GFP-labeled interneurons moved faster on the double knockout slice ( $20.8 \pm 0.6 \mu\text{m/h}$   $n = 4816$ , versus  $14.4 \pm 0.4 \mu\text{m/h}$   $n = 7016$ ,  $P < 0.0001$ , using unpaired  $t$ -test with Welch's correction). The mean and instantaneous velocities of the interneurons fit with previously published results (Tanaka et al., 2003).

These results suggested that the enhanced motility of the interneurons on the double KO substrate may result from a non-cell autonomous phenotype. One secreted protein, which has shown to affect migration of interneurons within this area is SDF-1 (stromal cell-derived factor-1) (Stumm et al., 2003). There was a clear difference in the pattern of *Sdf-1* expression in the cortex; in the double KO it extended from lateral to medial aspect of cortical anlage (Figure 8I) in comparison to the normal expression pattern seen in control brains (Figure 8J). Next, we examined the expression pattern of *Netrin-1*, which has been shown to affect the migration of interneurons (Stanco et al., 2009). We detected increased expression of *Netrin-1* in the striatum and lateral part of the cerebral cortex in comparison with the expression observed in control brains (Figures 8K,L). Collectively, our data indicate that the double KO mice exhibit earlier arrival of thalamocortical fibers, enhanced and faster migration of interneurons to the cortex, which is in part non-cell autonomous.



## DISCUSSION

Our study revealed an unexpected activity of the PAF-AH catalytic subunits acting as negative regulators of the Wnt pathway. In addition, deletion of the PAF-AH catalytic subunits increases the proliferation of GABAergic neurons in the ganglionic eminences and affects their migration. The advanced migration of GABAergic



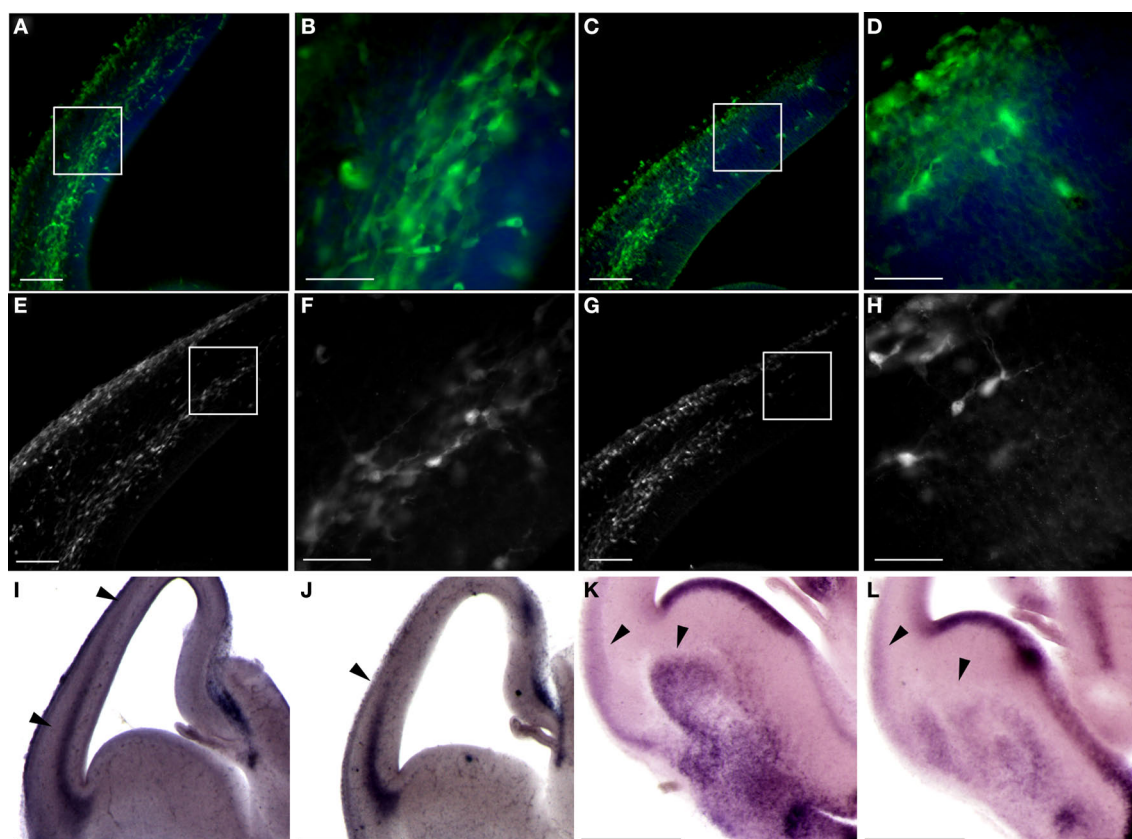
neurons is in part a non-cell autonomous effect which may be mediated through the modified expression of Wnts, SDF1 and Netrin1 in the double KO embryos.

## PAF-AH CATALYTIC SUBUNITS MODULATE THE Wnt PATHWAY

Increased expression of the PAF-AH subunits repressed gene expression mediated by TCF/LEF1 binding sites following activation with CA  $\beta$ -catenin or following activation of the pathway using Dvl. Subsequent to expression of PAF-AH catalytic subunits,  $\beta$ -catenin was mainly retained in the cytoplasm in a cultured cell line. Conversely, genetic elimination of the PAF-AH catalytic subunits resulted in an increase in the number of cells expressing nuclear  $\beta$ -catenin, increased proliferation in the GE and increased and expanded expression of upstream and downstream Wnt pathway components.

It may be of particular interest to repress the Wnt pathway, which is one of the five major signaling pathways predominating early and late embryonic development (Gerhart, 1999; Nusse,





**FIGURE 8 | Double KO mice exhibit advanced tangential migration.** (A–D) GAD67-GFP-labeled interneurons are found in more advanced positions in double KO brain sections (A,B) in comparison with control sections (C,D). Scale bars in (A) and (C) is 100  $\mu$ m, and in (B) and (D) is 40  $\mu$ m. (E,F) Calbindin immunostaining, which identifies a subgroup of interneurons, demonstrates advanced positioning of these interneurons in

the double KO (E,F) versus control (G,H). Scale bars in (E) and (G) is 100  $\mu$ m, and in (F) and (H) is 40  $\mu$ m. (I,J) *Sdf-1* expression by *in situ* hybridization is expanded in the cortex of double KO (I) versus the control cortex (J). (K,L) *In situ* hybridization of *Netrin-1* reveals an expanded expression in the striatum of double KO (K) versus the control striatum (L). Scale bars in (I–L) are 0.5 mm.

1999). The Wnt pathway participates in multiple roles as diverse as embryonic segmentation and patterning, gut patterning, nervous system development, formation and patterning of appendages, and stem cell proliferation. Most of the current information involves Wnt signaling through stabilization of  $\beta$ -catenin, which in turn enters the nucleus and controls gene expression (Nelson and Nusse, 2004). The Wnt pathway is highly conserved in evolution (Brown et al., 2008; Croce and McClay, 2008). We hypothesize that the evolutionary conserved function of the PAF-AH catalytic subunits is modulation of the Wnt pathway and not their interaction with LIS1 that acts as cytoplasmic dynein regulator (for review see Reiner, 2000). We are basing this hypothesis on several important observations. On one hand, *Lis1* heterozygote mice exhibit inhibition in radial and tangential migration (Hirotsune et al., 1998; Cahana et al., 2001; McManus et al., 2004b). On the other hand, PAF-AH catalytic subunit KO mice, do not exhibit a cortical phenotype, although they have decreased levels of the LIS1 protein (Koizumi et al., 2003; Yan et al., 2003). Dosage sensitivity of the LIS1 protein is well established. Reduced or increased LIS1 levels exhibit brain phenotypes in mice and in human (Hirotsune et al., 1998; Cahana et al., 2001; Reiner et al.,

2002; Bi et al., 2009). Therefore, the finding that PAF-AH catalytic subunit KO mice do not exhibit a cortical phenotype despite the apparent reduction in LIS1 levels was surprising. A possible explanation may be that the reduction in the LIS1 protein levels is just the portion of LIS1, which is required for interaction with the catalytic subunits, and the remaining amount of LIS1 required for other intracellular interactions is normal. An additional argument relates to the evolutionary conservation of the PAF-AH catalytic subunits. We have characterized the fly orthologs of the PAF-AH complex. The *Drosophila* alpha-subunit ortholog lacks two of the three active-site residues and is catalytically inactive against PAF-AH 1B substrates. The beta-subunit (LIS1) ortholog is highly conserved from *Drosophila* to mammals and is able to interact with the mammalian alpha-subunits but is unable to interact with the *Drosophila* alpha-subunit (Sheffield et al., 2000). It will be interesting to investigate the possibility that the *Drosophila* alpha-subunit can modulate the Wnt pathway.

Canonical Wnt signaling mediated by  $\beta$ -catenin has been proposed to function in both neural progenitor cell expansion and neuronal lineage choice (Chenn and Walsh, 2002; Chenn and Walsh, 2003; Zechner et al., 2003; Hirabayashi et al., 2004; Israsena et al.,

2004; Wrobel et al., 2007). The outcome of modulating the Wnt pathway is highly dependent on the brain area and the involvement of other factors (such as FGF2) (Israsena et al., 2004). Furthermore, some of the above mentioned studies used expression of stabilized  $\beta$ -catenin, which is not responsive to normal regulatory mechanisms. Within the medial GE, targeted deletion of  $\beta$ -catenin specifically impaired proliferation without grossly altering differentiation. The phenotype observed there was primarily due to loss of canonical Wnt signaling (Gulacsi and Anderson, 2008). Thus, it may be expected that increased Wnt signaling within the GE will result in increased proliferation as noticed in our studies.

#### PAF-AH CATALYTIC SUBUNITS AFFECT MIGRATION OF INTERNEURONS

We observed that the double KO mice exhibited increased proliferation in the GE, earlier development of thalamocortical fibers and enhanced and faster migration of interneurons to the cortex, which is in part non-cell autonomous. These phenotypes may be due to activation of the Wnt pathway, nevertheless, we cannot exclude the participation or crosstalk with additional signaling pathways. The phenotypes observed in the PAF-AH double KO mice resemble in part phenotypes of *Robo1*-deficient mice. *Robo1*-KO mice exhibited an increased number of interneurons that was due to increased proliferation. A similar increase in proliferation was observed in *Slit1(-/-)/Slit2(-/-)* mutant mice, suggesting that cell division was influenced by Slit-Robo signaling mechanisms (Andrews et al., 2008). In addition, the sonic hedgehog (Shh) pathway participates in proliferation regulation in the GE (Xu et al., 2005; Fuccillo et al., 2006; Gulacsi and Anderson, 2006). It will be interesting to examine the possible involvement of the Slit-Robo and Shh signaling in regulation of the phenotypes observed in the PAF-AH double KO mice.

Furthermore, in *Robo1*-deficient mice more interneurons migrated into the cortex (Andrews et al., 2006). It has been postulated that it was due to migration of the interneurons along the corticothalamic and thalamocortical axons that arrived earlier to their respective targets in the mutant than in the control (Andrews et al., 2006). In the PAF-AH double KO mice we observed earlier arrival of the thalamocortical axons identified by TAG-1 immunostaining. It is known that early born (E11.5–E14.5) GABAergic cells colocalize with TAG-1-positive axons *in vivo*, whereas later-born cells colocalized with TAG-1-negative fibers, suggesting temporal differences in substrate specificity (McManus et al., 2004a). Thus, for the embryonic day we examined (E13.5) TAG-1 should be a good marker for progression of GABAergic cells. In addition, we examined directly the position of the interneurons using GAD67-GFP knock-in mice. The GFP positive interneurons were found in more advanced positions in the PAF-AH double KO mice. When we transplanted wild-type GAD67-GFP interneurons on double KO or wild-type substrates we noticed faster motility on the mutant background. These results suggest a non-cell autonomous activity, which may be attributed to multiple known secreted factors. To date, multiple guidance molecules have been identified (for examples see Lopez-Bendito et al., 2008; Marin and Rubenstein,

2001). In our study, we examined three secreted factors; *Netrin-1*, *Sdf-1* and *Wnt7B*. *Netrin-1* was shown to interact with alpha3beta1 integrin to promote interneuronal migration (Stanco et al., 2009). However, *Netrin-1* activity is known to be context dependent and can act as either a chemoattractant or a chemorepellent (Alcantara et al., 2000; Charron et al., 2003; Wilson and Key, 2006). *Netrin-1* expression in the striatum acts as a repulsive factor (Hamasaki et al., 2001). Thus, increased expression of *Netrin-1* in the striatum of double KO may be involved in promoting interneuron migration to the cortex. In addition, *Netrin-1* expression may also influence thalamocortical projections (Powell et al., 2008). SDF-1 regulates migration and positioning of CXCR4-expressing interneurons during cortical development (Stumm et al., 2003; Liapi et al., 2008). Furthermore, localized expression of SDF-1 can induce chemotaxis in migrating neurons and alter their local trajectories (Liapi et al., 2008). The expanded and increased expression of *Sdf-1* in mice lacking PAF-AH catalytic subunits may explain in part the change in the migration rates observed in these mice. In addition, we noted increased and expanded expression of *Wnt7B*. To the best of our knowledge, there are no direct data suggesting for a role of this Wnt family member in regulation of interneuron migration. Nevertheless, the Wnt pathway participates in regulation of cell migration in a very diverse system which is the movement of groups of cells forming the lateral line primordium of the zebrafish (Aman and Piotrowski, 2008). In that system, Wnt/ $\beta$ -catenin works in conjunction with FGF signaling to regulate the expression of SDF-1 chemokine receptors. Thus, despite the long list of known secreted factors guiding and regulating interneuron migration, additional factors may be added. We suggest that possible modulation of the Wnt pathway by the PAF-AH catalytic subunits attenuates interneuron migration to the cerebral cortex.

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