Development/Plasticity/Repair

Retinoic Acid Induces Blood-Brain Barrier Development

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The blood– brain barrier (BBB) is crucial in the maintenance of a controlled environment within the brain to safeguard optimal neuronal function. The endothelial cells (ECs) of the BBB possess specific properties that restrict the entry of cells and metabolites into the CNS. The specialized BBB endothelial phenotype is induced during neurovascular development by surrounding cells of the CNS. However, the molecular differentiation of the BBB endothelium remains poorly understood. Retinoic acid (RA) plays a crucial role in the brain during embryogenesis. Because radial glial cells supply the brain with RA during the developmental cascade and associate closely with the developing vasculature, we hypothesize that RA is important for the induction of BBB properties in brain ECs. Analysis of human postmortem fetal brain tissue shows that the enzyme mainly responsible for RA synthesis, retinaldehyde dehydrogenase, is expressed by radial glial cells. In addition, the most important receptor for RA-driven signaling in the CNS, RA-receptor β (RAR β), is markedly expressed by the developing brain vasculature. Our findings have been further corroborated by *in vitro* experiments showing RA- and RAR β -dependent induction of different aspects of the brain EC barrier. Finally, pharmacologic inhibition of RAR activation during the differentiation of the murine BBB resulted in the leakage of a fluorescent tracer as well as serum proteins into the developing brain and reduced the expression levels of important BBB determinants. Together, our results point to an important role for RA in the induction of the BBB during human and mouse development.

Introduction

The vascular system of the CNS arises early in embryogenesis through the invasion of vascular plexus-forming angioblasts into the head region (Risau and Wolburg, 1990), followed by invasion of the CNS by vascular sprouts from the perineural vascular plexus, extending toward the ventricles (Greenberg and Jin, 2005). Peripheral vascular system development has been described in detail previously, and various signaling systems taking part in vasculogenesis, angiogenesis, and differentiation have been uncovered (Jain, 2003; Aird, 2007a,b). However, few reports exist on developmental CNS-specific cues for the induction of the specialized endothelial cell (EC) phenotype found at the blood—

presence of interendothelial tight junctions, transporters, and efflux systems, together ensuring a CNS environment that favors proper neuronal function (Pachter et al., 2003; Abbott et al., 2010).

brain barrier (BBB). This specialized EC phenotype includes the

CNS cells surrounding the endothelial layer are thought to provide angiogenic ECs with the appropriate signals for BBB maturation (Stewart and Wiley, 1981), as well as signals required for maintenance of the mature BBB phenotype. During CNS development, radial glial cells provide structural and trophic cues (Chanas-Sacre et al., 2000) and differentiated astrocyte end-feet projections provide an almost complete enveloping of the brain microvasculature in adult vertebrates (Mathiisen et al., 2010). The search for CNS-specific signals that affect the BBB phenotype in brain ECs has implicated astrocytes and glial progenitors as inducers of a specific BBB-phenotype in brain EC (Abbott et al., 2006).

Radial glial cell-derived retinoic acid (RA) is crucial in providing the brain with the correct developmental pattern during neurogenesis (Környei et al., 2007), and RA has complex and pleiotropic functions during vertebrate development (for review, see Niederreither and Dollé, 2008). However, the role of RA as a developmental cue in the CNS vasculature has remained uninvestigated. RA is a retinol-derived lipophilic metabolite, synthesized by retinaldehyde dehydrogenases (RALDH1–RALDH3), and one of the first described powerful morphogens during de-

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velopment. RA signaling has been implicated in cell fate determination for a wide range of cells in developing mammals. RA signaling can only occur when retinol, bound to retinol binding protein (RBP), is taken up from the circulation. Uptake of RBP-bound retinol occurs via the membrane-bound protein stimulated by retinoic acid gene 6 (STRA6) (Kawaguchi et al., 2007). Retinol is oxidized to retinaldehyde and further oxidized to retinaldehyde by RALDH enzymes, leading to freely diffusible RA, mainly acting in a paracrine manner. Target cells that express the nuclear RA receptors α , β , or γ (RAR $\alpha/\beta/\gamma$) and the scaffolding retinoid X receptors α , β , or, γ (RXR $\alpha/\beta/\gamma$) react to RA by the heterodimerization of RAR and RXR and subsequent transcription of genes with an upstream RA response element (RARE) (for review, see Duester, 2008).

In the current study, we set out to investigate the role of RA in BBB development. We provide insight in the spatiotemporal expression pattern of RALDH and RAR β during human CNS development and show that RA mediates various aspects of brain ECs, both *in vitro* and *in vivo*.

Materials and Methods

Cell culture. The human brain EC line hCMEC/D3 (Weksler et al., 2005) was provided by Dr. P.-O. Couraud (Institut Cochin, Université Paris Descartes, Paris, France) and grown in endothelial cell basal medium-2 supplemented with human EGF, hydrocortisone, GA-1000, FBS, VEGF, human fibroblast growth factor (FGF)-B, R3-IGF-1, ascorbic acid, and 2.5% fetal calf serum (Lonza). To obtain astrocyte cultures, fetal tissue (cerebral hemispheres) was obtained at 17–23 weeks of gestation following Canadian Institute of Health Research-approved guidelines. Astrocyte isolation has been described previously (Wosik et al., 2007). Human fetal astrocytes were used between P2 and P4, and cultures were determined to be >90% pure, as determined by glial fibrillary acidic protein (GFAP) immunostaining. Astrocyte-conditioned media (ACM) was harvested once a week from confluent flasks of human fetal astrocyte cultures.

Generation of recombinant hCMEC/D3 RARB knock-down cell lines. To establish knockdown of RARB, we used a vector-based shRNA technique. shRNA was purchased from Sigma. Recombinant lentiviruses were produced by cotransfecting subconfluent human embryonic kidney 293T (HEK 293T) cells with the shRNA lentivirus expression plasmid and packaging plasmids (pMDLg/pRRE and pRSV-Rev) using calcium phosphate as a transfection reagent. HEK 293T cells were cultured in DMEM supplemented with 10% FCS and 1% penicillin/streptomycin, in a 37°C incubator with 5% CO₂. The RARβ-specific shRNA sequence was 5'-CTGGGTAAATACACCACGAAT-3'; nontargeting shRNA control vector was bought from Sigma. Infectious lentiviruses were collected 48 h after transfection. The supernatant was centrifuged to remove cell debris and stored at -80°C. hCMEC/D3 cells were transduced with the lentivirus containing shRNA, and, after 48 h, cells stably expressing the shRNA containing construct were selected by puromycin-containing (0.2 \times 10⁻⁷g/ml) selection medium. After 48 h of selection, cells were maintained in normal medium.

Donor brain material. Immunohistochemistry was performed on tissue sections obtained from the brain collections of the Department of Neuropathology of the Academic Medical Center, University of Amsterdam (Amsterdam, The Netherlands). Parental consent was obtained for the use of brain tissue and for access to medical records for research purposes. The tissue was obtained from spontaneous or medically induced abortions, with appropriate parental written consent for brain autopsy. We also obtained normal-appearing control cortex at autopsy from two young cases (3 months and 7 months), without a history of seizures or other neurological diseases. All autopsies were performed within 12 h of death. Clinical data are presented in Table 1. Ten percent buffered formalin-fixed, paraffin-embedded tissue blocks were sectioned at 6 μ m and mounted on organosilane-coated slides (Sigma) for immunohistochemistry. Frozen tissue blocks were sectioned at 6 μ m and mounted on Superfrost slides (Microm).

Table 1. Clinical information regarding the brain tissue used in this study

Age	Sex	Brain weight	Postmortem delay (h)	Fixation time (weeks)	Reason for termination of pregnancy
10 gw	М	ND	ND	3	Spontaneous abortus
16 gw	F	15 g	7	3	Spontaneous abortus
20 gw	F	23 g	6	Frozen	Heart defect
22 gw	M	24 g	6	3	Spontaneous abortus
29 gw	M	131 g	8	3	Heart defect
3 months	M	572 g	7	3	
7 months	М	962 g	9	3	

ND, Not determined or available; gw, gestational week; M, male; F, female.

Animal experiments. To achieve specific disturbance of RA signaling in the developing embryo, pregnant C57BL/6 mice (Charles River Laboratories) were treated with validated in vivo RAR antagonist BMS493 [4-[(1E)-2-[5,6-Dihydro-5,5-dimethyl-8-(2-phenylethynyl)-2-naphthalenyl]ethenyl]benzoic acid] (Wendling et al., 2000) (5 mg/kg; Tocris Bioscience) or vehicle (DMSO) at 1:10 in nut oil. Hereto, mice were mated overnight, and the day of vaginal plug detection was marked as E0.5. Treatment was given by oral gavage twice daily, in 10-12 h intervals, and started at E10.5, during 6 consecutive days. At gestational day 16.5, the mice were killed, and the embryos were isolated from the uterus. To assess the permeability of the embryonic BBB, Cadaverine-Alexa Fluor 555 (5 mg/ml; Invitrogen) was injected intravenously 1 h before embryo isolation. Embryos were immediately fixed in 4% paraformaldehyde for 1 h to prevent diffusion of the tracer. For quantitative analysis of gene expression levels, embryonic brains were isolated and RNA isolation was performed using Trizol (Invitrogen) according to the protocol of the manufacturer. Downstream cDNA synthesis was performed as described below. For immunohistochemical analysis of protein expression, whole embryos were cryoprotected with 30% sucrose in PBS, embedded in 1.5% agarose, and snap frozen in liquid nitrogen. Whole embryos were cryosectioned as described above. Immunohistochemical analysis of tight junction protein zona occludens 1 (ZO-1), the endothelial marker platelet/endothelial cell adhesion molecule 1 (PECAM1), mouse IgG as an endogenous leakage marker, and neuronal nucleus marker (NeuN) are described under immunohistochemistry. The Animal Experiments Committee of the Free University Medical Center approved all of the experiments described in this study.

Immunohistochemistry. For single labeling of RAR β in human tissue sections, sections were deparaffinized and treated with 0.3% H_2O_2 in methanol for 20 min to reduce endogenous peroxidase activity. Antigen retrieval was achieved by incubating the sections at 100°C in citrate buffer (10 mm in PBS, pH 6.0) for 10 min. After washing with PBS, sections were treated with 0.1% saponin in PBS, washed, and subsequently incubated with primary antibody, polyclonal rabbit α RAR β (Abcam), in PBS overnight at 4°C. Slides were then washed and incubated with Envision+Dual link reagent (Dako) for 30 min, followed by visualization with the peroxidase substrate diaminobenzidine (Dako). After a short rinse in tap water, sections were incubated with hematoxylin for 1 min and extensively washed with tap water for 10 min. Finally, sections were dehydrated with ethanol, followed by xylene and mounted with Entellan (Merck).

Fluorescent immunohistochemistry for RAR β was performed using polyclonal rabbit α RAR β (Abcam) and monoclonal mouse α -PECAM1 (Dako). Tissue sections were stained as described above for single labeling; only peroxidase activity quenching was omitted, and all incubation steps were performed in the presence of 1% normal serum of the secondary antibody host (goat or donkey). Secondary antibodies used were goat α -mouse IgG–Alexa Fluor-488, donkey α -rabbit IgG–Alexa Fluor-555, and streptavidin–Alexa Fluor 488 (Invitrogen). After incubation with appropriate secondary antibodies, slides were washed and embedded in vinol mounting medium supplemented with 0.4% 4',6-diamidino-2-phenylindole (Invitrogen) to stain nuclei.

Immunohistochemistry on mouse embryonic sections was performed as described above with the following antibodies: rat α -mouse IgG biotinylated (Vector Laboratories); rat α -PECAM-1–Alexa Fluor-647 (Dako); rabbit α -NeuN (Millipore); and rabbit α -ZO-1 (Invitrogen). Fluorescence analysis was performed with a Leica DM6000 microscope

equipped with Leica advanced fluorescence software. Quantification of immunoreactivity was performed using NIH ImageJ software by setting a threshold value of fluorescence above the background and calculating the area above threshold per micrograph. At least five different micrographs per group were used in the analysis.

For RALDH staining in the developing human cortex, cryosections were air dried and fixed in acetone for 10 min. Sections were incubated as described above, overnight at 4°C with primary antibodies: mouse α -GFAP–Cy3 (Sigma); rabbit α RALDH1/RALDH2 (van de Pavert et al., 2009) (Abcam); and mouse α -vimentin, clone V9 (Millipore). After each incubation with a primary antibody, sections were washed and subsequently incubated with the appropriate secondary antibodies [donkey α -rabbit IgG–Alexa Fluor-488 and goat α -mouse IgG–Alexa Fluor 647 (Invitrogen)], washed, and embedded as described above. Confocal fluorescence analysis was performed with a Leica TCS SP2 confocal microscope equipped with Leica confocal software.

Immunocytochemistry. Fluorescent immunocytochemistry was performed as described for fluorescent immunohistochemistry, except for fixation method. Cultures of recombinant RAR β or scrambled shRNA hCMEC/D3s were grown to confluence in eight-well μ -slides (Ibidi), washed with ice-cold PBS, and fixed in 4% paraformaldehyde in PBS for 10 min at 4°C. Fixed cells were shortly permeabilized with 0.1% Triton X-100 in PBS, washed, and incubated overnight at 4°C with primary antibodies [mouse α -VE-cadherin (BD Biosciences) and rabbit α -ZO-1 (Invitrogen)], washed, and subsequently incubated with appropriate secondary antibodies: donkey α -rabbit IgG–Alexa Fluor-488 and goat α -mouse IgG–Alexa Fluor-647 (Invitrogen). Actin filaments were visualized with rhodamine phalloidin (Invitrogen).

Electric cell-substrate impedance sensing assay. ECISTM model 1600R (Applied BioPhysics) was used to measure the transendothelial electric resistance (TEER) of human hCMEC/D3 cell monolayers. A measure previously described to have a close inverse correlation with smallmolecule permeability (Santaguida et al., 2006). A total of 100,000 cells were seeded onto each well of an 8W10+ electric cell-substrate impedance sensing (ECIS) array (Ibidi) coated with collagen, and impedance was measured at multiple frequencies in real time. Cells were either treated with a 1:1 mixture of ACM or control medium and EGM-2 containing 2.5% FCS, with or without 5 µM RAR antagonists (Kagechika, 2002) LE135 (Tocris Bioscience) and LE540 (Wako Pure Chemicals) immediately after seeding. For ECIS measurements with an RAR agonist, cells were treated with indicated concentrations of RA (Sigma) or vehicle (100% EtOH) directly after seeding. For specific RAR β activation, the indicated concentrations of RAR β agonist and RAR α/γ antagonist BMS453 (Tocris Bioscience) or vehicle (100% DMSO) were added directly after seeding. BMS453 has confirmed activity in vitro (Germain et al., 2004) and in vivo (Matt et al., 2003). All ECIS measurements were furthermore subjected to a mathematical model to calculate the component of resistance attributed to cell-cell interactions, called barrier resistance (Rb), and the component derived from average membrane capacitance of cells covering the electrode (membrane capacitance) (Giaever and Keese, 1991).

Real-time quantitative PCR. Gene expression analysis was performed on confluent monolayers of hCMEC/D3 cells in 24-well microplates (Corning) treated with either 5 μ M RA for various lengths of time or confluent monolayers of recombinant hCMEC/D3 cell lines expressing either RAR β shRNA or scrambled shRNA. mRNA was isolated using an mRNA capture kit (Roche) according to the instructions of the manufacturer. cDNA was synthesized with the Reverse Transcription System kit (Promega) following the guidelines of the manufacturer, and RT-PCR was performed as described previously (García-Vallejo et al., 2004). Primer sequences used are as follows, describing the forward and reverse sequence, respectively: human: CDH5, TGACGTGAACGACAACTGGC and GACGCATTGAACAACCGATG; ABCB1, GTCCCAGGAGCCCATCCT and CCCGGCTGTTGTCTCCAT A; OCLN, CCCGTTTGGATAAAGAATTGG and TCAAACAACTTGGC ATGAGC; VEGF-A, CATCTTCAAGCCATCCTGTGTG and GCATGGTG ATGTTGGACTCCT; TJP-1, CCCGAAGGAGTTGAGCAGGAAATC and CCACAGGCTTCAGGAACTTGAGG;SLC2A1,GCAGCAGCCTGTGTAT GCC and AAGGCCGTCTTGACGATACC; RARB, GCAGAGCGTGTAAT TACCTTGAAandGTGAGATGCTAGGACTGTGCTCT;RARA,AGATTA

CTGACCTGCGAAGC and CCCTCTGAGTTCTCCAACATTTC; RARG, CCTTGTCTTTGCCTTTGCTG and CAGGTCCATGCGGTCTC; and GAPDH, CCATGTTCGTCATGGGTGTG and GGTGCTAAGCAGTTGG TGGTG; murine: Cdh5, CAGCAACTTCACCCTCATAAAC and TCCCGA TTAAACTGCCCATAC; Tjp-1, AGCGAATGTCTAAACCTGGG and TC CAACTTGAGCATACACAGG; Abcg2, GTTACCACTGTGAGCCCTA and TCTGTCTTGTTTGCTTCATGG; Rarb, GCCTGCAGAAGTGCTTTG AAGT and GCTCTCTGTGCATTCCTGCTTT; Gfap, CAGACTTTCTCC AACandCTCCTGCTTCGAGTC; Rbfox3, ATCCTTACCATCACACCATC G and GGAGACGGTGGAAGGTTTC; Pdgfrb, AGTGATGTCTGGTCT and TGGCATTGTAGAACT; and Pecam1, TGGTTGTCATTGGAGTGG TC and TTCTCGCTGTTGGAGTTCAG. Expression levels of transcripts obtained with real-time PCR were normalized to GAPDH expression levels. All conditions investigated were measured as quadruplicates.

Western blotting. Protein expression levels of VE-cadherin and ZO-1 were investigated in confluent monolayers of hCMEC/D3 cells grown in six-well microplates (Corning), treated with indicated concentrations of RA or vehicle for 48 h. Cells were washed with ice-cold PBS and lysed in cell lysis buffer (Cell Signaling Technology) containing complete protease inhibitor cocktail (Roche) at 4°C for 30 min. Cell lysates were then taken up in SDS sample buffer (100 mM Tris-HCl, pH 6.8), 4% SDS, 20% glycerol, and 5% β -mercaptoethanol) and heated to 95°C for 5 min. Lysates were then resolved on SDS-PAGE in triplicate, blotted, and incubated overnight with the primary antibodies mouse α -VEcadherin (Santa Cruz Biotechnology), rabbit α -ZO-1 (Invitrogen), and goat α-Actin (Santa Cruz Biotechnology) in Odyssey blocking buffer (LI-COR) diluted 1:1 in PBS, after initial blocking with blocking buffer for 1 h at room temperature. Primary antibodies were detected and quantified by incubation with appropriate IRDye secondary antibodies (at room temperature for 1 h in blocking buffer) and the Odyssey infrared imaging system (LI-COR). Actin quantification was used to correct for total protein loading.

Promoter-reporter constructs. The lentiviral vector pRRL-cPPT-CMV-X2-PRE-SIN (kindly provided by Dr. J. Seppen, Department of Experimental Hepatology, Academic Medical Center, Amsterdam, The Netherlands) was modified for use as a carrier of promoter-reporter cassettes as described previously (Fontijn et al., 2008). The reporter plasmid pGL-RARE-luciferase containing trimerized RAREs in front of the firefly luciferase gene was described previously (Weston et al., 2002). RARE promoter-reporter cassettes were transferred from pGLRAREluciferase as NotI-SalI fragments containing successively a synthetic pA/ transcriptional pause site, a RARE promoter fragment, the firefly luciferase reporter gene, and the simian virus 40 (late) polyadenylation signal. The resulting lentiviral RARE promoter-firefly luciferase reporter vector was then packaged, in the presence of a small amount (5% w/w) of the herpes simplex virus thymidine kinase Renilla luciferase lentiviral vector (Fontijn et al. 2008), using HEK 293T cells. Virus-containing supernatants were used for transductions, and luciferase activities were measured after at least 2 d, after 24 h of ACM exposure as described for the ECIS experiments using the dual luciferase system (Promega). Renilla luciferase levels were used to normalize firefly luciferase activity.

Statistical analyses. Data analysis was performed using GraphPad Prism software (version 5.01; GraphPad Software). Results are shown as mean \pm SEM, and statistical analysis was performed with either unpaired, two-tailed Student's t test or with one- or two-way ANOVA and subsequent Bonferroni's post hoc test correction. The applied test for each calculated value is described in the figure legends.

Results

Human fetal astrocytes increase brain endothelial barrier formation by a RA signaling pathway

To gain insight into RA production in the CNS, we first analyzed the expression of the RA synthesizing enzyme RALDH1/2 in developing and neonatal human brain material. Immunohistochemistry analysis of RALDH1/2 showed a distinct radial expression profile during human cortical development (Fig. 1A). Colocalization studies with antibodies against the radial glial cell markers vimentin and GFAP revealed ample expression of

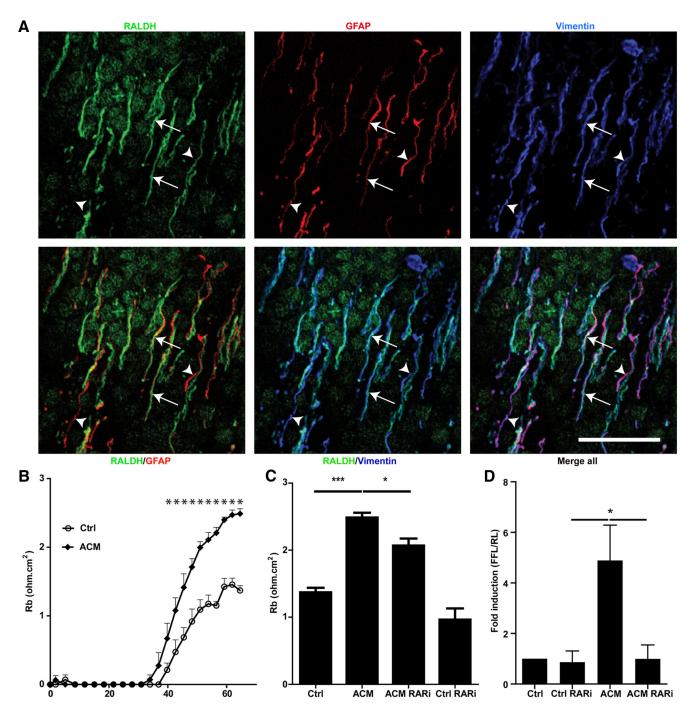


Figure 1. Fetal astrocyte-derived RA increases the resistance of brain ECs. *A*, Immunohistochemical analysis in human brain showing the expression of RALDH (green), GFAP (red), and vimentin (blue) in fetal cortex at 20 weeks of gestation. Note that RALDH-positive structures mostly overlap with both GFAP and vimentin-positive cells (arrows). RALDH-positive structures negative for GFAP and vimentin were also detected (arrowheads). Scale bar, 50 μm. *B*, Rb (0hms per square centimeters) of hCMEC/D3 cells during stimulation with fetal astrocyte factors is increased over time compared with control medium.* reflect significant differences compared with control (2-way ANOVA). *C*, Quantification of maximal Rb reached during an experiment similar to *B*. The addition of RAR antagonists LE135 and LE540 (ACM RARi) decreases the barrier-enhancing effect of fetal astrocyte factors on hCMEC/D3s (ACM), whereas vehicle-treated cells (Ctrl RARi) show no significant decrease compared with the control condition (Ctrl) (1-way ANOVA). *D*, Quantification of RARE–luciferase activity in hCMEC/D3 cells after ACM stimulation compared with control medium (Ctrl). The increase in luciferase activity is blocked by addition of RAR antagonists LE135 and LE540 (ACM + RARi). Addition of RAR antagonists to control medium-treated cells (Ctrl RARi) has no effect on luciferase activity (1-way ANOVA). *p < 0.05; ****p < 0.05; ****p < 0.001; n = 4 for all experimental conditions.

RALDH1/2 in radial glial cells during development. The presence of RALDH1/2 was not fully restricted to radial glial cells, suggestive of RA-synthesizing capacity for other cell types, such as neuronal precursors as well. To investigate the role of RA in the communication between fetal astrocytes and ECs of the BBB, we next established the effect of soluble fetal astrocyte factors on one of the hallmark characteristics of brain ECs, the formation of a

tight barrier. For this, hCMEC/D3s were seeded at confluence, and TEER was measured over time. Continuous exposure of the endothelial monolayers to ACM significantly increased TEER levels compared with unconditioned medium (maximum of $258.0 \pm 63.7 \Omega$; n = 4). To investigate the different components contributing to TEER, Rb was calculated and plotted over time. Fetal astrocyte factors also induced a significant increase in Rb

levels (maximum of $1.1 \pm 0.1 \Omega/\text{cm}^2$; n = 4; Fig. 1*B*), whereas cell spreading and proliferation were unaffected (data not shown). This indicates that the barrier-enhancing effect of fetal astrocyte factors is mainly attributable to increased function of interendothelial junctions.

To test our hypothesis that RA signaling is required for the increase in endothelial barrier function induced by ACM, we made use of the RAR antagonist combination LE135 and LE540 to block RAR activation by RA. The previously observed increase of Rb formation by fetal astrocyte factors was partially blocked by inhibition of the RARs in hCMEC/D3s (maximum of 37.34 ± 11.39% of increase; n = 4; Fig. 1C). Finally, to investigate whether exposure to fetal astrocyte factors leads to the activation of an RA signaling pathway in hCMEC/D3s, a RARE-luciferase reporter construct was introduced into hCMEC/D3s before exposure. Exposure to ACM resulted in an almost fivefold increase in endothelial luciferase expression compared with control medium (Fig. 1D). Notably, the ACM-induced increase in RARE-driven luciferase expression could be completely abolished by LE135/ LE540, showing that exposure to fetal ACM results in RARdependent RARE activation. Together, these results implicate RA signaling in the barrier-enhancing effect of fetal astrocyte factors on human brain ECs.

RA signaling in human brain ECs leads to increased barrier formation and the expression of BBB-specific genes

To gain more insight into the RA-mediated induction of BBB properties in hCMEC/D3s, we used various concentrations of all-*trans* RA (RA) in our *in vitro* ECIS assay. Continuous RA treatment resulted in a concentration-dependent increase of Rb of a confluent monolayer hCMEC/D3s over time when compared with vehicle-treated cells (Fig. 2A, maximal resistance (arrow): $0.1~\mu$ M, $9.8~\pm~1.9\%$ increase; $1~\mu$ M, $15.4~\pm~2.4\%$ increase; $5~\mu$ M, $21.0~\pm~2.5\%$ increase; n=4). Continuous treatment with $5~\mu$ M RA resulted in a 21% increase of the maximal Rb value reached in vehicle-treated cells, whereas cell spreading and proliferation were unaffected (data not shown).

The BBB phenotype is dependent on the expression of various proteins that contribute to tight junctions (VE-cadherin and occludin), transendothelial transport of nutrients [glucose transporter 1 (Glut-1)], and efflux pumps [P-glycoprotein (P-gp)]. To assess whether RA signaling in hCMEC/D3s regulates BBBrelated gene expression, we investigated the gene expression levels of P-gp, VE-cadherin, occludin, and Glut-1, after RA treatment. Treatment of confluent cultures of hCMEC/D3s with $5 \mu M$ RA for 48 h leads to increased gene expression levels of P-gp $(318.6 \pm 1.5\% \text{ of control})$, VE-cadherin $(172.2 \pm 24.9\% \text{ of con-}$ trol), and occludin (162.3 \pm 6.5% of control) (Fig. 2B). Glut-1 gene expression was only significantly increased after 6 h of RA stimulation (131.7 \pm 6.7% of control). Conversely, the gene expression level of the permeability-related factor VEGF-A is decreased during RA stimulation. As expected, RA treatment increased the expression of the known RA target gene RAR β $(283.2 \pm 35.7\% \text{ of control}).$

To determine protein expression of the RA-induced barrier genes, we performed Western blot analysis of two important determinants of BBB integrity, ZO-1 and VE-cadherin. ZO-1 and VE-cadherin protein expression was significantly and concentration dependently increased after treatment with either 1 or 5 μ M RA for 48 h compared with vehicle controls (ZO-1: 1 μ M RA, 218.2 \pm 49.6% of control; 5 μ M RA, 302.4 \pm 42.4% of control; VE-cadherin: 1 μ M RA, 114.8 \pm 2.5% of control; 5 μ M RA, 127.7 \pm 9.7% of control; Fig. 2C). The increase in protein expres-

sion for both VE-cadherin and ZO-1 was found to be functionally localized to cell–cell junctions by immunocytochemical analysis (Fig. 2D).

Together, these data show that RA is capable of inducing a tight barrier phenotype in human brain ECs by influencing gene and protein expression of molecules that are important for barrier function.

RAR β in human brain ECs is critical for barrier integrity

Because RAR β is the most important receptor in CNS-specific RA signaling (Mey, 2006; Wong et al., 2006), we used BMS453 as a specific RAR β agonist and RAR α/γ antagonist in our in vitro systems. In hCMEC/D3 cells, RARβ is the most abundantly expressed RAR subtype (Fig. 3A). Treatment of hCMEC/D3 cells with BMS453 resulted in a concentration-dependent increase in Rb over time compared with the vehicle control (Fig. 3B, maximal resistance (arrow): 0.1 μ M BMS453, 11.3 \pm 2.4% increase; 1 μ M BMS453, 16.7 ± 2.3% increase; 5 μ M BMS453, 27.2 ± 1.7% increase; n = 4). Moreover, RAR β activation resulted in a relative increase of the maximal Rb highly similar to comparable concentrations of RA (Fig. 2B). Cell spreading and proliferation were similar between all conditions (data not shown). To further enhance our understanding of the function of RAR β at the BBB, we specifically knocked down its expression in hCMEC/D3s by using a lentivirus encoding RAR β -specific or a nontargeting shRNA. Both recombinant cell lines were able to form a confluent monolayer under standard culture conditions. However, when compared with control cells, hCMEC/D3s with reduced RARβ gene expression (33.8 \pm 11.3% of control; Fig. 3E) failed to form normal Rb. Whereas RA still enhanced the barrier in control cells expressing nontargeting shRNA, RA stimulation was not able to restore the impaired Rb formation in RAR β -deficient cells (Fig. 3C). Together, the results revealed a role for constitutive RAR β activity in hCMEC/D3 barrier formation under normal culture conditions. To investigate the molecular alterations underlying the observed changes on Rb formation, we performed fluorescent immunocytochemistry analysis of two important barrier determining molecules, VE-cadherin and ZO-1, on confluent monolayers of both cell lines. Although actin distribution and expression was comparable between cell lines, a striking decrease of both VE-cadherin and ZO-1 immunostaining was observed in the cells with reduced RAR β expression compared with the control cells (Fig. 3D). Furthermore, the overlapping pattern of VEcadherin and ZO-1 is lost in the areas in which junctional expression of VE-cadherin or ZO-1 is preserved. The loss of a BBB phenotype after RAR $oldsymbol{eta}$ knockdown was also apparent from gene expression analysis of barrier-related genes (Fig. 3E). RARβ knockdown resulted in decreased gene expression levels of VEcadherin (25.4 \pm 8.1% of control), Glut-1 (42.2 \pm 4.2% of control), and ZO-1 (43.4 \pm 13.7% of control). However, P-gp mRNA levels remained unaltered by RAR β knockdown, suggesting a pathway other than RAR β activation in the RA-induced upregulation of P-gp gene expression. Finally, permeabilityrelated factor VEGF-A was highly increased during RARβ knockdown (211.3 \pm 14.4% of control). These experiments show that part of the barrier function of hCMEC/D3s is dependent on the presence of RAR β .

RAR β expression is located in microvascular structures during embryogenesis

Together, our experiments not only point out that RAR β is involved in RA-induced barrier function but also suggest a role for RAR β in the development of the BBB. To address this hypothesis,

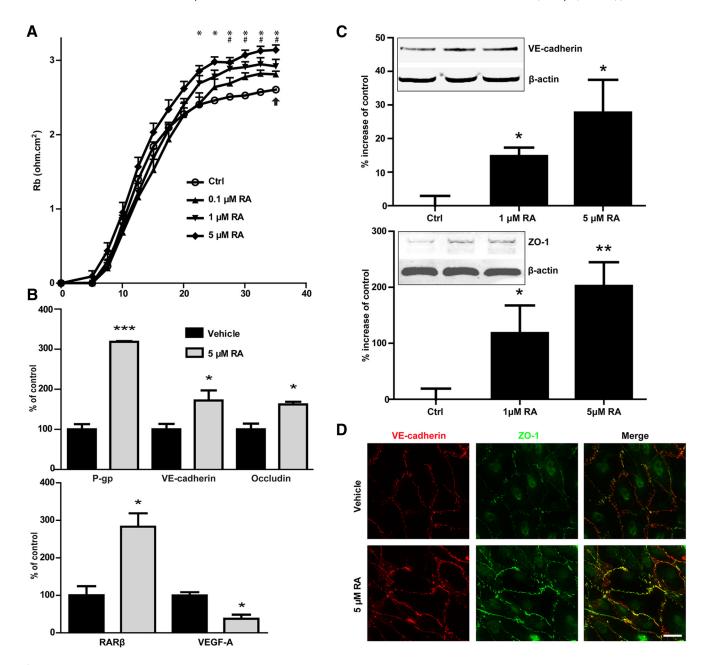


Figure 2. RA increases barrier formation of brain ECs by inducing barrier-related gene expression. **A**, Rb formation over time of hCMEC/D3s increases during RA treatment in a concentration-dependent manner compared with vehicle-treated cells (Ctrl). Arrow indicates the maximal Rb value. * and # reflect significant differences compared with control for 5 and 1 μ m RA, respectively (2-way ANOVA). **B**, Expression levels of barrier-related genes are increased after RA (48 h, 5 μ m) treatment. Gene expression level of permeability-related VEGF-A is decreased, whereas gene expression level of RA-target gene RAR β is increased, after RA (48 h, 5 μ m) exposure (2-tailed Student's *t* test). **C**, Quantification of VE-cadherin and Z0-1 protein expression as measured by Western blot, corrected for actin. Both Z0-1 and VE-cadherin protein levels in whole-cell lysates of hCMEC/D3s stimulated with RA (48 h, 5 μ m) increase significantly in a concentration-dependent manner (1-way ANOVA). Insets show representative images of the specific bands measured, both experiments performed in triplicate. **D**, Immunocytochemical analysis of VE-cadherin (red) and Z0-1 (green) expression of hCMECD3 cells stimulated as described in **C**. Note that the RA-induced increase in expression of both proteins localizes to the cell–cell junctions. Scale bar, 20 μ m. *p < 0.05; **p < 0.001.

we analyzed RAR β expression in human fetal brain tissue, in various stages of development, by immunohistochemistry (Fig. 4A). As early as 10 weeks of gestation, RAR β expression is evident in structures with vascular morphology but can also be found in multiple other cell types, and many structures with vascular morphology are devoid of RAR β expression. Vascular RAR β expression becomes more pronounced at the later developmental stages of 22 and 29 weeks of gestation when most vascular structures are RAR β positive. Moreover, its expression coincided with the BBB markers ZO-1 and occludin. Glut-1 immunoreactivity precedes the presence of RAR β , with pronounced vascular immunoreac-

tivity from 10 weeks of gestation. Costaining with an antibody against PECAM1 confirms the vascular nature of RAR β -positive cells (Fig. 4B). Assessment of RAR β expression in 3 (data not shown) and 7 (Fig. 4A) months postnatal human cortical tissue revealed no abundant vascular expression pattern. Whereas the expression pattern of ZO-1, occludin, and Glut-1 remain present in postnatal tissues, RAR β expression is mainly restricted to neurons. A similar developmental pattern can be observed by analyzing RAR β gene expression in whole mice brain lysates from different developmental stages, corrected for vascular content (PECAM1). Relative RAR β expression shows a decline as the

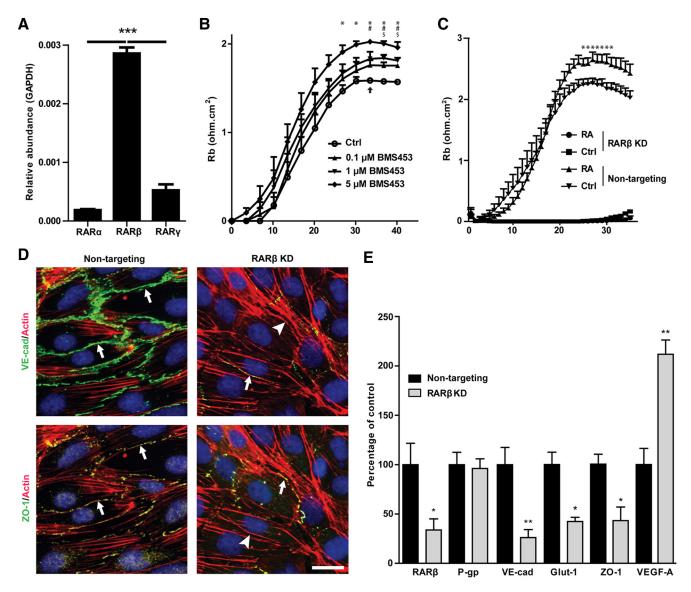


Figure 3. RAR β is the dominant RAR involved in barrier formation. *A*, Relative abundance of RAR subtype transcripts in hCMEC/D3 cells compared with GAPDH (1-way ANOVA). *B*, Rb formation of hCMEC/D3 over time rises with increasing concentrations of RAR β agonist BMS453. Arrow indicates the maximal Rb value reached. *, #, and \$ reflect significant differences compared with control for 5, 1, and 0.1 μ m, respectively (2-way ANOVA). *C*, Rb values of hCMEC/D3 cells transduced with nontargeting shRNA are increased by stimulation with 5 μ m RA. *reflect significant differences of nontargeting + RA compared with control (2-way ANOVA). *D*, Immunohistochemical analysis showing anti-VE-cadherin or ZO-1 (green) and actin fibers (red). hCMEC/D3 cells transduced with nontargeting shRNA (nontargeting) show junctional and overlapping localization of VE-cadherin and ZO-1 (arrows), as well as normal actin distribution. hCMEC/D3 cells transduced with RAR β shRNA (RAR β KD) show highly decreased VE-cadherin and ZO-1 staining. Junctional localization of VE-cadherin and ZO-1 (arrows) and actin distribution is preserved, although little to no overlap between VE-cadherin and ZO-1 is observed (arrowheads). Scale bar, 20 μ m. *E*, Barrier-related gene expression levels, as well as RAR β gene expression level, are decreased in RAR β shRNA-expressing hCMEC/D3 cells compared with nontargeting shRNA. However, P-gp expression level remains unaffected by the knockdown of RAR β gene expression, whereas the gene expression level of VEGF-A is highly increased (2-tailed Student's *t* test). *p < 0.05; *p < 0.001; *

brain develops and is significantly lower in adult brain compared with all earlier stages (Fig. 4C). The presence of RAR β in vascular structures during human brain development and the simultaneous appearance of BBB markers ZO-1 and occludin suggest a role for RAR β in the maturation process of brain ECs during a specific time window early in development, resulting in the loss of vascular RAR β expression shortly before or after birth.

RA signaling is necessary for BBB formation in the developing mouse brain

To investigate the relevance of RA signaling in BBB development *in vivo*, we treated pregnant mice with the RAR antagonist BMS493, from gestational age 10.5 to 16.5. By starting at 10.5 d of gestation, we can exclude disturbance of normal brain develop-

ment until the nervous vascular plexus starts to invade the CNS parenchyma (Walls et al., 2008). BMS493 treatment resulted in a distinct embryonic appearance, including smaller size, incomplete eye development and pigmentation, and an edematous formation in the neck, corresponding with that found in RA-deficient animals (Quadro et al., 2005). The permeability of the embryonic cerebral vessels (PECAM1 positive) was assessed by immunohistochemical detection of endogenous IgG and the accumulation of circulating Cadaverine—Alexa Fluor-555. In the CNS of embryos from vehicle-treated pregnant mice, IgG immunoreactivity was exclusively located within the vasculature, whereas Cadaverine—Alexa Fluor-555 was not detected. However, the CNS of embryos from BMS493-treated pregnant mice displayed a distinct extravascular presence of both IgG and Ca-

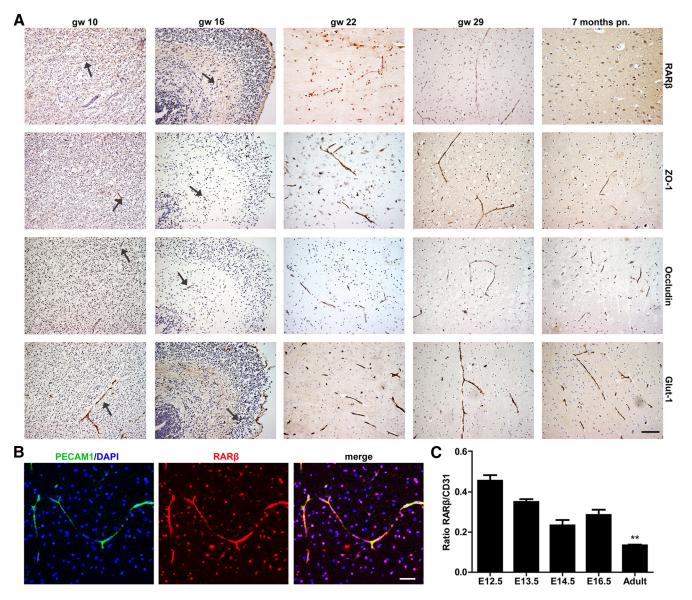


Figure 4. RAR β expression in the neurovasculature during cortical development. **A**, Immunohistochemical analysis of RAR β , Z0-1, occludin, and Glut-1 in paraffin sections of human cortical brain tissue at various stages of development. Note that from 10 weeks of gestation (gw), RAR β is present in vascular structures (arrows). The vascular staining pattern becomes more pronounced in later stages of development (22 and 29 weeks of gestation) and is absent in postnatal brain [7 months postnatal (pn.)]. Z0-1 and occludin immunoreactivity profiles follow that of RAR β (arrows), whereas Glut-1 immunoreactivity in vascular structures precedes vascular RAR β immunoreactivity. Scale bar, 50 μm. **B**, Immunohistochemical analysis showing RAR β (red) and PECAM1 (green) at 29 weeks of gestation; note that the PECAM1-positive structures overlap with RAR β , showing both nuclear and cytoplasmic RAR β presence in fetal microvessels. Scale bar, 20 μm. **C**, Gene expression levels of RAR β in mouse whole-brain lysates during different developmental stages, corrected for PECAM1 gene expression levels (1-way ANOVA). ***p < 0.01; p = 4.

daverine-Alexa Fluor-555 in cells surrounding the PECAM1positive vasculature. Double-labeling experiments with the neuronal nuclear antigen NeuN showed that both the IgG and Cadaverine-Alexa Fluor-555-positive cells are neurons (Fig. 5*A*). The average amount of IgG immunoreactivity and Cadaverine-Alexa Fluor-555 fluorescence per micrograph was found to be significantly higher in BMS493-treated embryonic brain sections compared with DMSO-treated embryos (IgG, 357.8 ± 103.5% of control, p = 0.037; Cadaverine–Alexa Fluor-555, 1650 \pm 554.5% of control; p = 0.025, two-tailed Student's t test). To analyze the BBB-related gene expression, the embryonic brains were isolated to study CNS gene expression levels only. BMS493-treated embryonic brains showed a significant decrease in gene expression levels of the major RA target gene RAR β compared with vehicletreated controls ($-57.0 \pm 10.9\%$ of control; Fig. 5B). This confirms BMS493 efficacy to reduce RAR activation in the brain. Importantly, BMS493 treatment did not affect the density of brain capillaries or PECAM1 gene expression levels, nor did it affect the expression of other brain cell-specific genes for astrocytes (GFAP), neurons (NeuN), and pericytes (PDGFRB) (data not shown). The expression levels of BBB-related genes ZO-1 $(-32.2 \pm 8.6\% \text{ of control})$, VE-cadherin $(-40.1 \pm 14.2\% \text{ of })$ control), and the ABC transporter gene breast cancer resistance protein (BCRP; $-34.9 \pm 14.2\%$ of control) were significantly reduced in BMS493-treated embryos compared with vehicletreated embryos (Fig. 5B). Fluorescent immunohistochemical analysis of ZO-1 protein expression in BMS493-treated embryos revealed that ZO-1 is localized to PECAM1-positive structures but clearly showed a reduced immunoreactivity when compared with vehicle-treated embryo brain (Fig. 5C). BMS493 treatment did not significantly affect claudin-5 and occludin gene expression levels. Together, these data show that RA and RARs are

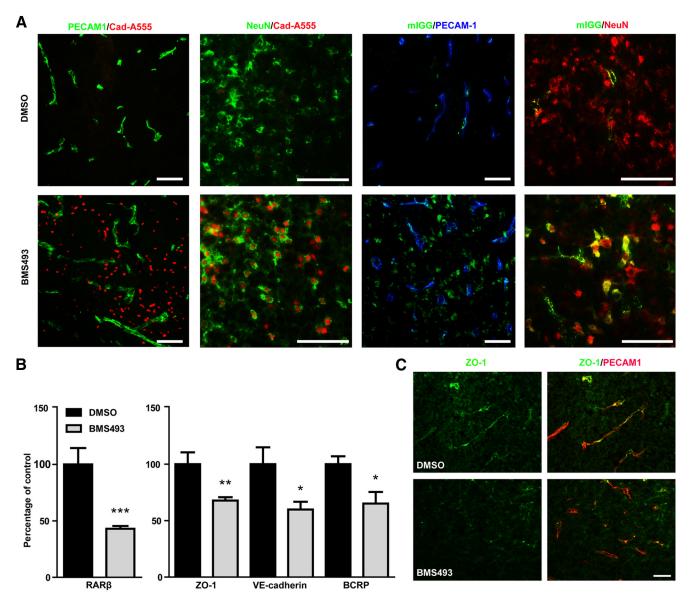


Figure 5. RAR antagonist BMS493 administration in pregnant mice results in increased permeability of the embryonic BBB, coinciding with decreased BBB gene expression. **A**, Immunohistochemical analysis showing endogenous IgG (mIGG, green) or Cadaverine—Alexa Fluor-555 (Cad-A555, red) with PECAM-1, and NeuN in embryonic brain of vehicle- and BMS493-treated embryos. Note that both the extravascular IgG-positive cells and Cadaverine—Alexa Fluor-555-positive cells have NeuN-positive nuclei/cytoplasm. Scale bars, 50 μ m. **B**, RAR β gene expression level and gene expression levels of barrier-related genes are decreased in BMS493-treated embryonic brain compared with vehicle controls (2-tailed Student's t test). **C**, Immunohistochemical analysis showing Z0-1 (green) and PECAM1 (red). BMS493-treated embryonic brain shows a reduced immunoreactivity for Z0-1 with a junctional localization compared with vehicle-treated embryonic brain (arrowhead). Scale bar, 50 μ m. *p < 0.05; ***p < 0.01; ***p < 0.001; *

implicated in the development of the murine BBB and that a reduction of RAR activity in the CNS leads to structural BBB impairment.

Discussion

To date, this is the first investigation of RA signaling focusing on the development of the BBB. The present study shows that RA signaling is involved in the induction of BBB development. During fetal brain development, RA is synthesized by RALDH1/2-positive radial glial cells, and its corresponding receptor, RAR β , is markedly expressed in developing human brain microvasculature. Moreover, *in vitro* analyses revealed that RA increases the formation of a tight barrier and mediates the expression of BBB-related genes, including VE-cadherin, P-gp, and ZO-1, and conversely downregulated the permeability factor VEGF-A. Specific activation and knockdown studies identified RAR β as the major

RAR responsible for its barrier-enhancing effect in human brain ECs. Finally, treatment with an RAR antagonist caused a reduction in RAR activation during BBB development in mice and a decrease of BBB-related gene expression, functionally illustrated by the permeability to circulating IgG and exogenous tracer Cadaverine—Alexa Fluor-555.

RA is best known as a neuronal differentiation factor in the developing CNS. Interestingly, the developmental window in which the vasculature invades the CNS and matures into the BBB overlaps with the induction and patterning of neuronal differentiation and outgrowth (Tam and Watts, 2010), and both systems share certain guidance cues and differentiation-inducing signaling events during their development (Larrivée et al., 2009). Our findings point to RA as one of these common denominators of neuronal and vascular development.

Specific differentiation of BBB ECs is crucial in safeguarding the CNS environment and normal neuronal function. We here report that the astrocyte precursors, radial glial cells, which are in close proximity to the developing CNS vasculature, are capable of RA production through the expression of RALDH. The current view on BBB development relies heavily on signals from the resident CNS cells, such as astrocytes, microglia (Abbott et al., 2010), and, more recently, pericytes (Armulik et al., 2010; Daneman et al., 2010). However, underlying mechanisms are just now beginning to emerge in the form of CNS-specific Wnt/ β -catenin signaling (Liebner et al., 2008; Stenman et al., 2008; Daneman et al., 2009), Sonic hedgehog (Shh) signaling (Alvarez et al., 2011), and intracellular stabilization signals mediated by VE-cadherin (Rudini et al., 2008; Taddei et al., 2008). We propose that RA signaling acts in concert with the signaling profile of CNS-specific Wnt and Shh pathways during maturation of the BBB. Besides directly acting on the ECs of the developing BBB, RA has been described to act as a regulator of Shh and Wnt pathways in the developing CNS (Halilagic et al., 2007), thereby linking the three pathways and possibly providing a time window of action for each separate pathway. Furthermore, a recent report describes a regulatory role for RA on the major developmental pathways of FGF and Notch signaling during spinal cord development (Paschaki et al., 2012), of which the Notch pathway has already been linked to BBB development and integrity (Li et al., 2011).

The dependence of neural progenitors on radial glial cell-derived RA has been described previously (Környei et al., 2007; Kane et al., 2008; Wang et al., 2011). Moreover, the main receptor for RBP-bound retinol, STRA6, is expressed in the developing bovine CNS by a subset of large blood vessels and astrocytes surrounding the mainly STRA6-negative microvasculature (Kawaguchi et al., 2007). This implies that the ECs of the CNS microcirculation rely on RA synthesis by proximal radial glia. Another source of RA synthesis during CNS development was described in the form of neural progenitor cells (Engberg et al., 2010). This suggests that the role of RA synthesis might shift from one cell type to another as CNS development progresses. This is further strengthened by the finding that postnatal and adult rat astrocytes retain their capacity for RA synthesis, inducible by specific conditions (Shearer et al., 2012).

In vitro studies using a human brain endothelial barrier model revealed a potential working mechanism for RA at the developing BBB. Cultured fetal astrocytes induced an RA-dependent signal in brain ECs that lead to the modulation of different BBB aspects, including the formation of a tight barrier and the expression of BBB-specific proteins. The maturation of angiogenic vasculature into a functional BBB requires the formation of high paracellular resistance. We show that RA leads to increased expression of VE-cadherin, occludin, and ZO-1, three molecules described to play a critical role in barrier integrity (Abbott et al., 2010). Moreover, the endothelial BBB phenotype encompasses the polarized expression of specific transport and efflux proteins, forming the metabolic barrier. RA highly increases the gene expression of P-gp and to a lesser extent the expression of Glut-1, showing that RA regulates various aspects of the brain endothelial barrier. Previous reports on RA-induced gene expression in brain ECs are in line with our findings, showing increased expression of P-gp (El Hafny et al., 1997) and the detoxifying enzyme γ -glutamyl transpeptidase (Lechardeur et al., 1995).

The induction of BBB aspects in cultured brain ECs by RA was dependent on the expression of RAR β , which shows high relative abundance compared with the two other RAR subtypes and was mimicked by specific RAR β activation. Moreover, the antagonis-

tic effect of BMS453 on RAR α and RAR γ did not result in adverse effects on Rb value, showing that a functional barrier increase is not dependent on the activation of RAR α or RAR γ . The knockdown of RAR β expression resulted in loss of endothelial barrier function and reduced expression of BBB markers. P-gp expression was unaffected by RAR β knockdown, indicating that part of the metabolic barrier function might be regulated by RAR α/γ / RXR heterodimers or other nuclear receptors. Furthermore, previous reports have shown RAR-mediated repression of gene transcription in the absence of RA as a ligand (for review, see Niederreither and Dollé, 2008), which may underlie the observed changes in endothelial gene expression. The importance of RAR β at the BBB was further supported by expression analysis of the developing human brain. During fetal development of the CNS, RAR β is expressed in the ECs of the developing vasculature. This suggests their responsiveness to the local release of RA, and RARB expression is furthermore a clear indication of RA-dependent gene transcription (Liao et al., 2005). Vascular RAR β expression seems to be conserved between mice and humans, because RAR β gene expression decreases during mouse brain development when corrected for vascular content.

To translate our findings to in vivo BBB development, we pharmacologically blocked the activation of RARs in utero. A major drawback in animal models in which RA production is fully ablated is that exogenous RA is needed to ensure successful conception and embryonic CNS development. Moreover, animals with an RAR-subtype deficiency often show redundancy effects, limiting the usefulness of the phenotypical observations (Ghyselinck et al., 1997), with additional complications arising from the loss of normal transcriptional repression by RARs in the absence of RA. Because BBB development starts from E10, with the ingrowth of ECs from the neural vascular plexus (Walls et al., 2008), this time point was chosen to start BMS493 treatment. Analysis of the accumulation of an injected tracer, Cadaverine, as well as of endogenous IgG in embryonic brains at E16.5 revealed a distinct extravascular staining pattern in BMS493-treated animals. This was not observed in control embryos, suggestive of serum leakage across a structurally impaired BBB. Intensely stained extravascular IgG deposits and Cadaverine-positive nuclei were found in NeuN-positive cells surrounding the CNS microvasculature, suggesting neuronal uptake of extravasated IgG and Cadaverine. For IgG, the finding is in line with a study describing BBB damage in a model for temporal lobe epilepsy (Rigau et al., 2007), in which IgG-positive neurons were observed in the hippocampus after BBB damage, coinciding with a decrease of vascular ZO-1 expression. Comparable findings were described in an animal model for amyotrophic lateral sclerosis (ALS) (Alexianu et al., 2001), as well as in ALS patients (Engelhardt and Appel, 1990). The accumulation of Cadaverine-Alexa Fluor-555 in neurons has been described in mice after BBB dysfunction as a result of a lack of pericytes (Armulik et al., 2010) and apolipoprotein E4 deficiency (Bell et al., 2012). Accompanying the observed permeability to serum proteins, we found decreased expression levels of ZO-1, VE-cadherin, and BCRP, whereas the total number of PECAM1-positive vascular structures was unaltered. Furthermore, ZO-1 immunoreactivity on the microvasculature as well as a clear junctional localization was reduced in BMS493-treated embryonic brains. Considering the regulatory role for RA in BBB-associated pathways like Shh- and Wntsignaling, indirect effects of RAR-inhibition via these pathways might be considered. However, our in vitro studies describe a direct effect of RA on brain ECs and RAR β expression is located at the developing BBB in the human fetal CNS. Together, these

findings suggest a direct role for RA on endothelial maturation in the developing BBB.

The search for developmental cues involved in BBB formation has implications for neurodegenerative disorders in which BBB disruption is paramount. Many reports show the association of BBB breakdown with pathology of CNS disorders, such as multiple sclerosis (van Horssen et al., 2007; Kooij et al., 2010, 2011; Neuwelt et al., 2011), in which restoring BBB function may provide new ways to counteract these disorders. Developmental pathways activated by RA that induce BBB function could prove to be important targets to boost the self-regenerative capacity of the adult brain.

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