

Maternal dietary choline deficiency alters angiogenesis in fetal mouse hippocampus

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We examined whether maternal dietary choline modulates angiogenesis in fetal brain. Pregnant C57BL/6 mice were fed either a choline-deficient (CD), control (CT), or choline-supplemented diet (CS) from days 12 to 17 (E12-17) of pregnancy and then fetal brains were studied. In CD fetal hippocampus, proliferation of endothelial cells (EC) was decreased by 32% ($p < 0.01$ vs. CT or CS) while differentiated EC clusters (expressing factor VIII related antigen (RA)) increased by 25% ($p < 0.01$ vs. CT or CS). These changes were associated with >25% decrease in the number of blood vessels in CD fetal hippocampus ($p < 0.01$ vs. CT and CS), with no change in total cross-sectional area of these blood vessels. Expression of genes for the angiogenic signals derived from both endothelial and neuronal progenitor cells (NPC) was increased in CD fetal hippocampus VEGF C (*Vegfc*), 2.0-fold, $p < 0.01$ vs. CT and angiopoietin 2 (*Angpt2*), 2.1-fold, ($p < 0.01$ vs. CT). Similar increased expression was observed in NPC isolated from E14 fetal mouse brains and exposed to low (5 μ M), CT (70 μ M), or high choline (280 μ M) media for 72 h (low choline caused a 9.7-fold increase in relative gene expression of *Vegfc* ($p < 0.001$ vs. CT and high) and a 3.4-fold increase in expression of *Angpt2*, ($p < 0.05$ vs. CT and high). ANGPT2 protein was increased 42.2% ($p < 0.01$). Cytosine-phosphate-guanine dinucleotide islands in the proximity of the promoter areas of *Vegfc* and *Angpt2* were hypomethylated in low choline NPC compared to CT NPC ($p < 0.01$). We conclude that maternal dietary choline intake alters angiogenesis in the developing fetal hippocampus.

brain | VEGF | ANGPT2 | endothelial cell | diet

Choline is an essential nutrient for mammalian cells (1), and it is needed for membrane formation, for methylation, and for acetylcholine biosynthesis (2). Pregnant women in the United States do not eat diets delivering the recommended intake of choline (3). Large amounts of choline are delivered to the fetus across the placenta (4), resulting in very high tissue choline concentrations in the fetus and newborn (5, 6). Maternal choline deficiency during pregnancy alters neurogenesis in fetal mouse hippocampus (7, 8), and this effect is likely mediated by alterations in gene-specific DNA methylation within neuronal progenitor cells (NPC) that will form the hippocampus (9, 10). Rodent dams fed choline-deficient (CD) diets during late pregnancy had offspring with diminished progenitor cell proliferation and increased apoptosis in fetal hippocampus (11, 12), accelerated differentiation of NPC (13–15), insensitivity to long-term potentiation (LTP) when they were adult animals (16), and decremented visuospatial and auditory memory as adults (17–20). In utero development of the nervous system requires coordination of neurogenesis with angiogenesis to assure that neurons are supplied with oxygen and that waste products are removed (21–23). Therefore, maternal dietary choline also might modulate angiogenesis in fetal brain.

The balance between angiogenesis and neurogenesis is carefully modulated by local cues (growth factors, extracellular matrix) in the mesoderm (21, 24–27). Endothelial cells (EC), derived from mesodermal cells, generate blood islands early in gestation (embryonic day 6.5; E6.5) and by E9–10 the primordial vasculature is formed (21, 28). Angiogenesis and vascular maturation are

synergistically regulated by vascular endothelial growth factors and their receptors and by angiopoietin (ANGPT)/endothelial receptor tyrosine kinase (Tie-2) signaling (29–31). EC in the perineural plexus of a ten-day-old mouse embryo express high levels of VEGF-receptor 1 (VEGF-R1), VEGF-receptor 2 (VEGF-R2), and tyrosine kinase with immunoglobulin-like and EGF-like domains 1 (TIE-1) and TIE-2, while their ligands are expressed in the surrounding tissues. This pattern of growth factors stimulates the invasion of neuroectoderm (21). VEGF synthesis by neurons and by NPC lining the ventricular zone of the hippocampus stimulates capillary ingrowth towards this region (22, 32). Also, binding of VEGF to its receptors activates the antiapoptotic kinase Akt PI3K that inhibits activation of caspase 9 and caspase 3, thereby increasing endothelial cell survival (23). Finally, VEGF mediates axonal guidance (23) and neural progenitor cell migration (33). Upon VEGF-R2 activation, this receptor is phosphorylated, activating ras, p38MAPK, and ERK, directly resulting in increased cell proliferation and migration (23). Vascular maturation and stabilization are required for functional angiogenesis (34) and this is mediated by ANGPT (35, 36).

We examined whether EC proliferation and differentiation in fetal hippocampus were affected by maternal dietary choline, and we determined whether NPC, when exposed to low choline, expressed angiogenic signals.

Results

Maternal Choline Deficiency Decreased Maternal Hepatic Phosphocholine. Varying dietary choline intake for 5 d in pregnant dams reduced hepatic concentrations of phosphocholine as expected: CD, 201 nmol/g \pm 26 SE ($p < 0.05$ different from control (CT)); CT, 464 nmol/g \pm 51; and choline-supplemented (CS), 648 nmol/g \pm 74 ($n = 10$ /group). These findings match our previously published data on choline metabolites and maternal choline diet (8).

Choline Deficiency Increased mRNA Expression Encoding Angiogenic Signaling Molecules in Fetal Brain and in Cultured Neural Progenitor Cells. As discussed earlier, angiogenesis and vascular maturation are regulated by VEGF and its receptors as well as by ANGPT (29–31). Previous studies on gene expression profiling of NPCs indicated possible alterations in the mRNA levels of angiogenic related genes when exposed to choline deficiency (37). Progenitor cells isolated from E14 fetal mouse brains and grown in low choline (5 μ M) medium as compared to CT medium (70 μ M choline) had increased gene expression (relative to TATA box binding protein (*Tbp*) housekeeping gene) of *Vegfc* (9.7-fold, $p < 0.001$) and angiopoietin 2 (*Angpt2*) (3.4-fold, $p < 0.05$ vs. CT) (Fig. 1). The expression of these two genes also was increased in E17 fetal

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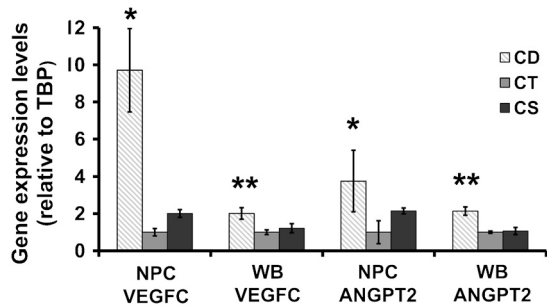


Fig. 1. Choline deficiency increases expression of *Vegfc* and *Angpt2* in cultured neural progenitor cells and in E17 fetal brain. E14 mouse brain neural progenitor cells were grown in low choline (5 μ M; CD), (70 μ M; CT), or (280 μ M; CS) medium for 72 h. Pregnant mice were fed a low, normal, or supplemented choline diet between days 12 to 17 of gestation and fetal hippocampi were collected at E17 ($n = 5$ pups from 5 dams/group). mRNA was isolated from NPC and whole brain (WB) and expression levels of *Vegfc* and *Angpt2*, relative to *Tbp* housekeeping gene, were assessed with qRT-PCR using the $2^{-\Delta\Delta CT}$ method. $n = 5$ plates/group. Data are presented as mean \pm SE. * $= p < 0.05$; ** $= p < 0.01$ different from CT and CS by REST analysis.

brain from CD as compared to CT and CS dams (*Vegfc* by 2-fold, $p < 0.01$; *Angpt2* by 2.1-fold, $p < 0.01$).

Choline Deficiency Altered the Protein Levels of ANGPT2 and VEGFC. We measured ANGPT2 protein levels in NPC samples exposed to low, normal, and high choline levels by ELISA. Choline deficiency increased ANGPT2 protein levels by 42.2% ($p < 0.01$ vs. CT) (Fig. S1B). A similar pattern of changes was observed for VEGFC (Fig. S1A).

Maternal Choline Deficiency Decreased Endothelial Cell Proliferation in the Fetal Hippocampus. The proliferation index (ratio of BrdU-Isolectin cells/total DAPI-isolectin cells) in EC was decreased in the dentate gyrus of the hippocampus in the CD group relative to the CT ($p < 0.0001$) or CS group ($p < 0.001$). A similar decrease was observed when the proliferation index in whole hippocampus was analyzed in CD and compared to CT ($p < 0.01$) or supplemented groups ($p < 0.01$) (Fig. 2 A, B; Fig. 3). There were no differences between groups in the surface of the dentate gyrus or whole hippocampus (inset Fig. 3).

Maternal Choline Deficiency Decreased the Number of Blood Vessels Formed in the Fetal Hippocampus. We explored the effect of decreased EC proliferation on capillary number and cross-sectional area. The blood vessels were counted per dentate gyrus and per whole hippocampus. The number of isolectin-labeled blood vessels in fetal hippocampus was decreased by >25% in CD as compared to CT or CS ($p < 0.01$) (Table 1). There were no significant changes in the average cross-sectional area of individual blood vessels or in the combined cross-sectional area of all the microvessels from one hippocampus.

Though we observed changes in the hippocampus region, there were no differences observed in the cingulate and midposterior neocortex (Table S1). We suggest that choline acts by changing epigenetic marks and that this may occur in progenitor cells only at a specific period when they are dividing and differentiating. Neurogenesis and angiogenesis in the cortical areas start before embryonic day 12, the beginning of our dietary manipulation. The regional specificity likely reflects these differences.

Maternal Choline Deficiency Increased the Number of Endothelial Cells Expressing Von Willebrand factor-Factor VIII Complex in the Fetal Hippocampus. As they differentiate, endothelial cell clusters synthesize Von Willebrand (VW) factor and Factor VIII in the Weibel-Palade bodies within the cells (38). The number of EC clusters

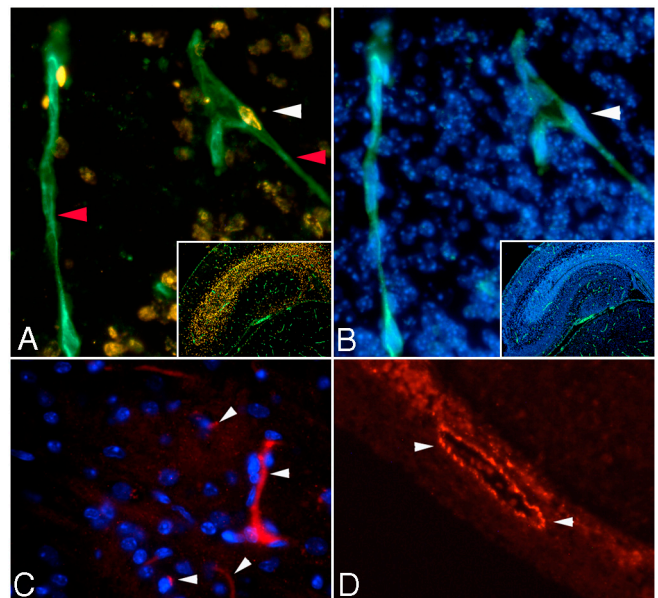


Fig. 2. Immunohistochemistry studies in fetal brain. A and B: Representative 200X fluorescent images of the CORNUS AMMONI region 1 (CA1) of E17 hippocampus. A multiplex immunolabeling was performed for BrdU (golden staining—ALEXA 555, white arrowhead A), isolectin IB4 (green fluorescence—ALEXA 488, red arrowhead A) and DAPI (blue fluorescence when bound to genomic DNA, white arrowhead B). The proliferation index of EC was calculated as the ratio between the number of BrdU positive nuclei colocalized with isolectin (A) and DAPI positive nuclei colocalized with isolectin (B). Small insets are images of the entire hippocampus area captured with 50X magnification (A—isolectin-BrdU and B—DAPI-isolectin). C and D: Factor VIII RA staining of E17 fetal mouse hippocampus. Representative 200X fluorescent images of blood vessels expressing Factor VIII RA in a 5 μ m hippocampal sections; image acquisition was performed with a rhodamine filter—blood cells fluoresce red (ALEXA 546 nm dye; white arrowheads) and nuclei fluoresce blue (DAPI staining). C shows small blood vessels with different trajectories and levels of Factor VIII as well as DAPI positive EC nuclei. D presents a more intense, Factor VIII positive, blood vessel with a bigger caliber situated in close proximity to the dentate gyrus.

expressing VW factor in fetal hippocampus was increased by 25% in the CD group as compared to CT and CS ($p < 0.01$) (Fig. 2 C, D, Fig. 4). A similar pattern was observed in the CA2, CA3 regions and dentate gyrus of the fetal hippocampus.

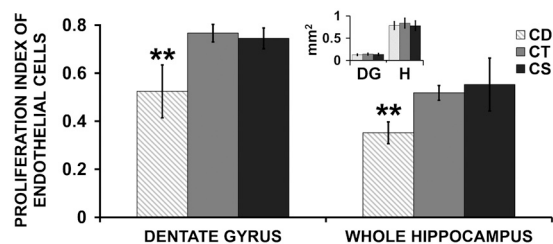


Fig. 3. Maternal choline deficiency decreases endothelial cell proliferation in E17 fetal hippocampus: Pregnant mice were fed CD, CT, or CS diet between days 12 to 17 of gestation and fetal brains were collected at E17. At E15, all dams received an intraperitoneal (i.p.) dose of BrdU. Brain sections were probed with isolectin IB4 and anti-BrdU antibodies. The proliferation index of EC was calculated as the ratio of colabeled BrdU-isolectin positive cells to DAPI-isolectin positive cells in both hippocampi (or both dentate gyri) of each brain in six consecutive 5 μ m sections. The surface area of whole hippocampus (H) and dentate gyrus (DG) were measured for each sample and found to be constant between the treatments (small inset). Data are presented as mean \pm SE. $n = 6$ pups from 6 dams/group. Groups were compared using one way ANOVA followed by Tukey-Kramer test. ** $= p < 0.01$

Table 1. Maternal dietary choline deficiency decreases the number of blood vessels in fetal hippocampus

Angiometric analysis	CD	CT	CS
Number of blood vessels per hippocampus	342 ± 19**	473 ± 21	455 ± 26
Combined cross-sectional area (μm ²)	29,507 ± 3,638	38,631 ± 3,734	39,193 ± 1,421
Individual microvessel area (μm ²)	101 ± 8	106 ± 7	121 ± 9

Pregnant mouse dams were fed a CD, CT, or CS diet from E12 to E17 when angiogenesis in fetal hippocampi was evaluated. Blood vasculature was fluorescently labeled using isolectin-ALEXA488 conjugated (green fluorescence) as described in *Materials and Methods* section. The number, total surface area, and the diameter of microvessels that were isolectin G5-IB4 positive were measured using the ImageJ Software in both hippocampi of one fetal brain and the average numbers were reported. Six consecutive sections (5 μm) of medial hippocampal region from each sample (30 μm total) were analyzed. Statistical analysis was performed by ANOVA and Tukey-Kramer. Data is presented as mean ± SE, *n* = 6 pups from 6 different dams/group, ** = *p* < 0.01 different from CT and CS groups.

Choline Deficiency Altered Methylation of Several CpG (Cytosine-Phosphate-Guanine Dinucleotide) Islands Downstream of the Transcription Starting Site (TSS) of *Vegfc* and *Angpt2*. Cytosine methylation modulates gene expression (39). We examined the methylation status of the first three CpG islands from *Vegfc* and *Angpt2* because of their close proximity to the TSS for each gene. Decreased methylated DNA in both *Vegfc* and *Angpt2* was detected (despite low signal compared to the positive CT (*Xist*)) in the low choline NPC for all islands (*p* < 0.01 vs. CT) (Fig. 5).

Discussion

We show, using a rodent model, that maternal dietary choline intake modulates angiogenesis in fetal brain. Maternal diets low in choline were associated with diminished proliferation of EC in fetal hippocampus. We suggest the following hypotheses: Choline is an important methyl-donor (9) and a diet low in choline in the pregnant dam results in decreased DNA methylation in fetal brain within the promoter of two genes that are important regulators of angiogenesis (*Vegfc* and *Angpt2*). Hypomethylation of genes is usually associated with increased gene expression (9), and *Vegfc* and *Angpt2* are over expressed in fetal brain. This change increases VEGFC and ANGPT2 angiogenic signaling, causing EC to differentiate more rapidly (and to secrete the proteins VW factor and Factor VIII). EC differentiation ends cell proliferation and this results in decreased numbers of blood vessels in the hippocampus. Previously published studies established that maternal dietary intake of choline during sensitive periods of neurodevelopment in the fetus altered hippocampal neurogenesis (8); in our current study, we learned that these effects are not limited to neuronal cells but extend to endothelial cells and blood vessel formation in the fetal hippocampus.

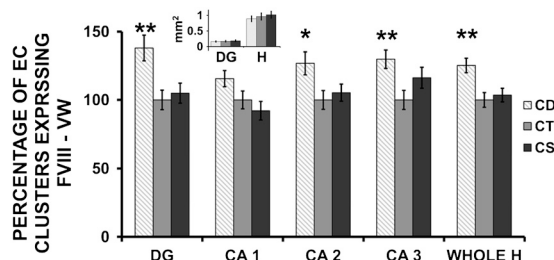


Fig. 4. Maternal choline deficiency increases the number of endothelial cells expressing Factor VIII and Von Willebrand factor in E17 fetal hippocampus. Pregnant mice were fed CD, CT, or CS diet between days 12 to 17 of gestation and fetal brains were collected at E17. Consecutive fetal brain sections were immunolabeled with anti-VW Factor VIII antibodies conjugated with ALEXA 546 nm. Clusters of endothelial cells expressing VW Factor were counted in E17 hippocampal sections (DG = dentate gyrus; CA1 = cornu ammonis 1; CA2 = cornu ammonis 2; CA3 = cornu ammonis 3; and Whole H = whole hippocampus). The surface area of whole hippocampus (H) and dentate gyrus (DG) were measured for each sample and found to be constant between the treatments (small inset). Data are presented as mean ± SE. *n* = 7 pups from 7 dams/group. * = *p* < 0.05, ** = *p* < 0.01, as compared to CT and CS groups by ANOVA with Tukey-Kramer tests.

As discussed earlier, angiogenesis and vascular maturation are regulated by VEGF and ANGPT-TIE2. Though CD was associated with increased *VEGFC*, it was paradoxically associated with decreased, not increased endothelial cell proliferation. This modification occurred because differentiation of EC was accelerated, and differentiated cells divide less frequently, if at all. We assessed endothelial cell differentiation using several proteins (and glycoproteins) that are specifically made by mature endothelial cells. One such protein is von Willebrand factor (vWf) which is stored in the Weibel-Palade bodies (25). This protein stabilizes Factor VIII, a protein involved in the clotting cascade (40). Expression of VW factor is higher in mature EC, and is minimal in early endothelial progenitor cells (38). This expression can be influenced by a number of factors including epigenetic mechanisms (41), hormones (17 β-estradiol) (42), and homocysteine (43) (note that choline availability is an important modulator of homocysteine (44)).

Modifications in the proliferation of EC can be associated with structural differences in the vascular tree that are reflected in capillary morphometry and number of blood vessels (21). The number of blood vessels in fetal brain was decreased in CD, but the total cross-sectional area was not significantly altered. This modification may be explained by increased levels of VEGFR ligands in the CD group, as this signal encourages formation of wide lumina with thin walls (21, 45). Studies on pupillary membrane capillaries show that increased ANGPT2 in the absence of VEGF increases EC death. Disruption of VE-cadherin junctions by ANGPT2 increases the response to VEGF stimulation in EC as a survival mechanism (46). ANGPT2 potentiates the effects of VEGF and therefore, the combination of VEGF and ANGPT2 promotes a rapid increase in capillary diameter and remodeling of basal lamina (46). VEGF also stimulates nitric oxide synthase (eNOS) in EC (47). Both eNOS and VEGF can promote microvasculature maturation in combination with increased ANGPT2 levels (48). Indeed, alterations in the choline levels induce high calretinin expression (15), a known marker of differentiation. VEGFC binds to VEGF-R2 and 3 and enhances lymphangiogenesis, angiogenesis, and neurogenesis (49). By activating VEGF-R2, VEGFC stimulates endothelial cell migration via P38MAPK and focal adhesion kinase (FAK) (23). It is possible that NPC lining the periventricular areas secrete VEGFC and ANGPT2 as survival factors (46) that also happen to stimulate angiogenesis.

Choline is an essential nutrient and a major source of methyl groups used to synthesize *S*-adenosylmethionine. Low choline levels alter global and gene-specific DNA and histone methylation (9, 10, 50). Therefore, we suggest that the gene expression changes in *Vegfc* and *Angpt2* are the result of epigenetic modifications. We analyzed the gene sequences of *Vegfc* and *Angpt2* and found several CpG islands downstream from the promoter area of these genes, but also at the TSS. These CpGs were hypomethylated in the low choline group; epigenetic modifications occurring near the promoter area, sometimes many kilobases from the TSS can alter gene expression by modulating the binding of enhancers/insulators to specific loci (51, 52). DNA hypomethylation decreases the recruitment of methyl binding domain (MBD)

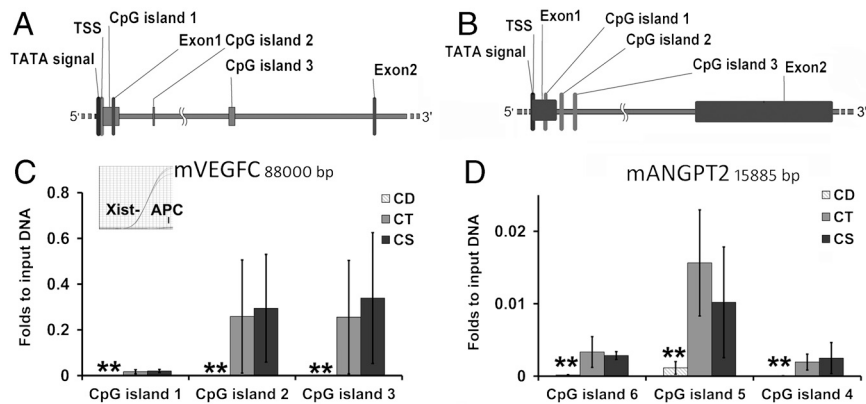


Fig. 5. Choline deficiency alters the methylation status of CpG islands within *Vegfc* and *Angpt2*. E14 NPC were grown in low, control of CS medium as described in Fig. legend 1 ($n = 4$ /group). Enzymatically sheared DNA samples were enriched in methylated cytosine following the MIRA protocol and analyzed by real time PCR as described in *Materials and Methods* section. *A* and *C*: Diagram (to scale) of *Vegfc* (80 Kb) and *Angpt2* (16 Kb), respectively, containing the first three CpG islands analyzed and the gene features in close proximity to these CpGs: exons 1 and 2, TSS. *B*: Methylation (folds to input DNA) of CpG islands within *Vegfc*. Inset shows the amplification curves of positive control *Xist* gene and no amplification for the negative control *APC* gene detected by real time PCR; *D*: Methylation (folds to input DNA) of CpG islands within *Angpt2*. Values are reported as mean \pm SE. Statistic analysis was performed with REST (rest pair wise reallocation randomisation test). ** = $p < 0.01$.

families of proteins such as MeCP2. Once bound to methylated CpG sites, MeCP2 silences transcription of downstream genes by virtue of its interaction with a histone deacetylase (HDAC)/Sin3 complex (53). This mechanism modifies chromatin and alters binding of transcription factors (53).

Computational analysis of CpG islands in *Vegfc* and *Angpt2* (Fig. S2) revealed several CpGs located near putative binding sites for a number of transcription factors (for *Vegfc*: *AP2*, *Sp1*, *PAX4*, *bHLH*, *NF-kB*, and *NRSF*; for *Angpt2*: *HMG*, *bHLH*, *ETS1*, *ARNT* and *PPAR*). Epigenetic modulation of transcription factor binding at one or all of these sites could alter angiogenesis. For example, Sp1 and AP2 enhancers increase *Vegf* expression during developmental angiogenesis and this is critical for vasculature formation (54, 55). In pathological situations, *Vegfc* and *d* expression can be modulated by NF-Kappa B activation (56). There is a NRSF binding site located in the third CpG island of the *Vegfc* gene; modifications of epigenetic marks at the NRSF binding site in *Calb1* gene in fetal brain were associated with altered gene expression (50). Similar mechanisms could alter VEGF signaling. *Angpt2* is up-regulated in endothelial cells by a hypoxia induced factor binding site located in its first intron and by Ets-1 (57). SOX18 is a member of the Sry-related HMG box-containing family of transcription factors and is expressed transiently in endothelial cells during the development of blood vessels. Mutations resulting in expression of dominant negative SOX18 severely impair vascular development (58). DNA bending induced by the HMG domain can facilitate the formation of higher-order nucleoprotein complexes altering H3K4 (59) methylation, suggesting that HMG domain proteins may have an architectural role in assembling such complexes (60). The aryl hydrocarbon receptor nuclear translocator (ARNT) serves as the obligate heterodimeric partner for bHLH-PAS proteins involved in sensing and coordinating transcriptional responses to hypoxia and developmental pathways (61). Lastly, PPARs upregulate angiogenic factors in other model systems (62).

The effects of maternal dietary intake of choline on angiogenesis in fetal brain could have great importance for humans. Only 14% of pregnant women in the United States consume adequate amounts of choline in their diets (and less than 5% of 18–30 young women who are not pregnant consume the recommended intake) (3). Approximately half of young women have a single nucleotide polymorphism in the *PEMT* gene and inefficiently form new choline moiety and therefore must eat choline or they develop organ dysfunction (63–65). These women are likely to need more choline during pregnancy to sustain normal fetal development

(66, 67). In the United States, pregnant women in the lowest quartile for choline intake have a 4-fold increased risk of having a baby with a birth defect compared with women in the highest quartile for choline intake (68, 69). Thus, humans eat marginal amounts of choline and a large portion of the population has increased dietary requirement for choline, making the effects of choline on fetal brain angiogenesis of potential public health importance.

Materials and Methods

Additional information in *SI Text*

Animals. Timed-pregnant C57BL/6 mice were used in all experiments according to a protocol described elsewhere (8). At the end of E11 they were randomly assigned to one of three feeding groups: CD (CD; AIN-76A diet with no choline), CT, or CS (CS; AIN-76A diet with 4.95 g/kg choline chloride) and fed these diets until they were killed on E17. On E15, a single intraperitoneal injection of 50 mg/kg body weight 5-bromo-2'-deoxyuridine (BrdU; Sigma-Aldrich, Inc.) was administered to the pregnant dams.

Fetal Brain Collection. On E17 the fetal brains from two male pups from each litter were collected using methods we previously published (8, 70). 5 μ m hippocampal sections were cut in coronal orientation. The remaining brains were frozen in liquid nitrogen and used for gene expression determinations.

Fetal Mouse NPC in Culture. E14 NPC from C57BL/6 mice were obtained from Lonza and plated according to the manufacturer's protocol. Small tertiary neurospheres were generated as previously described (50) and suspended in custom Neurobasal medium (D7005A, Atlanta Biologicals) containing 5 μ M choline chloride (CD), 70 μ M choline chloride (CT) or 280 μ M choline chloride (CS) with the ingredients previously described (50).

Immunohistochemistry in Paraffin Embedded Sections. To assess EC proliferation, a double immunolabeling technique using both BrdU (S-phase marker) (71) and isolectin IB4 (vasculature marker) (28, 72) was used. Developmental angiogenesis occurs in waves (73); therefore we evaluated EC proliferation over a period of 48 h (E15–E17). Similarly, equivalent sections were probed for Von Willebrand factor (Factor VIII related antigen, VW factor)—Factor VIII complex. In all cases, 4', 6-diamidino-2-phenylindole (DAPI, Sigma) 0.1 μ g/mL for 20 min was used for staining the nuclear DNA (full details provided in *SI Text*).

Image Analysis. Fluorescent images were collected with a Nikon FXA microscope (Nikon) equipped with an Optronics TEC-470 CCD Video Camera System (Optronics Engineering). The number of blood vessels and the average cross-sectional area of isolectin positive vessels were counted using the ImageJ Software version 1.37v (<http://rsb.info.nih.gov/ij>, NIH) with an integrated macro for analyzing particles. The proliferation index of EC was calculated as the ratio of colabeled BrdU-isolectin positive cells to DAPI-isolectin

positive cells in both hippocampi of each brain in six consecutive 5 μm sections. For analysis of capillary numbers and cross-sectional volume, we used consecutive sections totaling 30 μm of thickness from the medial-frontal region of the hippocampus from six fetuses from different dams. To assess the number of EC clusters that were VW Factor positive, the fluorescent structures were counted per entire hippocampus as well as for selected hippocampal regions: CA1 (Cornu Ammoni region 1), CA2, CA3 and DG (Dentate Gyrus).

Real-Time RT-PCR. RNA was extracted from cultured NPC or from whole brain samples using the RNeasy Mini Kit (Qiagen) transcribed to cDNA and 2 ng per PCR was amplified in the second step. TBP mRNA levels do not vary with choline availability in the central nervous system, as shown in published microarray studies (37). Primers were designed for *Angpt2* and for *Vegfc* and synthesized as described in *SI Text*. The conditions for the amplification reactions were: 95 °C for 10 min; 40 cycles of 94 °C for 15 sec, 56 °C for 30 sec, and 72 °C for 50 sec. The relative quantification method ($2^{-\Delta\Delta\text{CT}}$) was used to analyze gene expression changes as described elsewhere (74).

Methylated CpG Island Recovery Assay (MIRA). Genomic DNA was isolated from NPC samples using Qiagen DNaseasy minicolumns (Qiagen) following manufacturer's protocol (75), quantified with a Nanodrop 8000 spectrophotometer (Nanodrop) and stored in elution buffer at -80°C . In preparation for the enrichment step, 4,000 ng of gDNA from each sample were digested with *MseI* restriction enzyme for 2 h at 37 °C followed by heat-inactivation. Input DNA aliquots were incubated with recombinant His-MBD2b/MBD3L1 protein

complex followed by capture with protein G coated magnetic beads. *Vegfc* and *Angpt2* genomic sequences were retrieved from the NIH mouse genome database (<http://www.ncbi.nlm.nih.gov/nucore>) including the gene features (CpG islands and exons/introns locations). For each CpG island several primer pairs were designed and synthesized, and PCR amplification performed as described in detail in *SI Text*.

Maternal Hepatic Phosphocholine. Phosphocholine is the labile storage form for choline in liver and is an excellent indicator of dietary choline status (76). It was assayed using liquid chromatography—electrospray ionization— isotope dilution mass spectrometry as previously described (77).

Statistical Analysis. For gene expression the analysis was performed using the Relative Expression Software Tool—Multiple Condition Solver REST-MCS version 2 (free from the website <http://www.gene-quantification.de/rest.html>) which uses a pair wise fixed reallocation randomization test. All statistical analyses of epitope levels and vascular tree determinations were performed using JMP software (V 2; SAS Institute, Cary, NC) with ANOVA and Tukey-Kramer tests. Data are presented as mean \pm SE.

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