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Development of spatial integration depends on top-down and interhemispheric connections that can be perturbed in migraine: a DCM analysis

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Abstract In humans, spatial integration develops slowly, continuing through childhood into adolescence. On the assumption that this protracted course depends on the formation of networks with slowly developing top-down connections, we compared effective connectivity in the visual cortex between 13 children (age 7-13) and 14 adults (age 21–42) using a passive perceptual task. The subjects were scanned while viewing bilateral gratings, which either obeyed Gestalt grouping rules [colinear gratings (CG)] or violated them [non-colinear gratings (NG)]. The regions of interest for dynamic causal modeling were determined from activations in functional MRI contrasts stimuli > background and CG > NG. They were symmetrically located in V1 and V3v areas of both hemispheres. We studied a common model, which contained reciprocal intrinsic and modulatory connections between these

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E. Fornari (☒) · M. G. Knyazeva Department of Radiology, Centre Hospitalier Universitaire Vaudois (CHUV), and University of Lausanne, Rue du Bugnon, 46, 1011 Lausanne, Switzerland e-mail: Eleonora.Fornari@chuv.ch

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R. Rytsar · M. G. Knyazeva Department of Clinical Neuroscience, Centre Hospitalier Universitaire Vaudois (CHUV), and University of Lausanne, Rue du Bugnon, 46, 1011 Lausanne, Switzerland e-mail: Romana.Rytsar@gmail.com regions. An analysis of effective connectivity showed that top-down modulatory effects generated at an extrastriate level and interhemispheric modulatory effects between primary visual areas (all inhibitory) are significantly weaker in children than in adults, suggesting that the formation of feedback and interhemispheric effective connections continues into adolescence. These results are consistent with a model in which spatial integration at an extrastriate level results in top-down messages to the primary visual areas, where they are supplemented by lateral (interhemispheric) messages, making perceptual encoding more efficient and less redundant. Abnormal formation of top-down inhibitory connections can lead to the reduction of habituation observed in migraine patients.

Keywords Children · Effective connectivity · Inhibition · Predictive coding · Visual cortex · Functional magnetic resonance imaging

Introduction

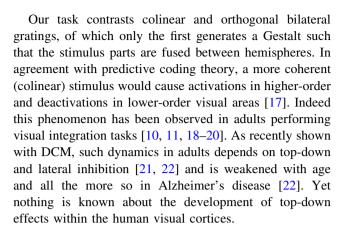
The developing brain provides a unique opportunity to investigate its functional networks in consecutive temporal windows. Due to the interregional heterochronicity of maturation, the role of long-developing functional circuits and their dysfunction in pathology can be understood. The top-down influences are based on such circuits with protracted development [1, 2]. Their extended development through adolescence has been shown for prefrontal top-down influences underlying voluntary behavioral control [3, 4]. However, top-down effects are involved in all types of cognitive activities, including perception, where they reshape interactions between distributed networks under different perceptual contexts.



Moreover, their deviant or delayed development might be suspected in the pathological conditions that affect distributed brain regions and have age-dependent prevalence. For instance, migraine, which is a frequent cause of headache in children, can be among such neurological disorders. Indeed, as MRI-based studies in adults show distributed cerebral networks are changed in chronic migraine patients [5, 6]. The prevalence of migraine increases from 3 to 19 % in children between 5 and 12-year-old, and then drops to some extent [7], suggestive of a link to certain protracted developmental processes.

The predictive coding hypothesis [8] persuasively offers a framework for the role of top-down effects in the hierarchical processing of sensory stimuli. This hypothesis suggests that at each level of signal processing, neural networks form internal models of the natural world based on its regularities. These representations generate predictions of how the sensory evidence should be interpreted and send them via descending connections upstream, where the model information is compared to the stimulus-driven activity. At the lower processing level, the well-predicted activity is removed by inhibition, and the residual error between the prediction and sensory evidence is calculated. The signal deviations from the predictions are sent to the higher processing centers for adjusting the model. Importantly, by inhibiting the predictable and, therefore, excessive activity, such a hierarchical mechanism optimizes processing of a stimulus.

Top-down influences can be studied non-invasively in terms of effective connectivity, which evaluates the influence that one local neural system (source) exerts on another (target) by means of dynamic causal modeling (DCM) [9]. For the analysis of feedback effects, it is important that DCM differentiates positive and negative couplings. Positive modulation expresses functional excitation in a target region in the case of activity increase correlated with the source, while negative modulation manifests functional inhibition in the target in the case of source activation. In this study, we examine the top-down and interhemispheric modulatory connections in a visual network, using a perceptual task that involves interhemispheric spatial integration [10, 11]. Spatial integration refers to the brain processes that implement a global representation of spatially extended objects by assembling local information across the visual field. In contrast to the majority of visual functions that achieve adult levels within a few months from birth, spatial integration develops at least until midadolescence [12–15]. Migraine interferes with the development of spatial integration such that, compared to agematched controls, children with migraine show reduced improvement in the performance of contour integration task between 6 and 14 years [16].



We hypothesized that the functional circuits providing top-down effects continue to advance during the second decade of human life. In addition to characterizing their developmental changes from a network perspective, we aimed at linking them to structural brain maturation as manifested by interhemispheric callosal connectivity. Here, we report the DCM-based evidence of this process in healthy children, a necessary step for the evaluation of the state of long-developing networks in children with neurological disorders including migraine. Some of these results have been mentioned elsewhere in relation to the corpus callosum development [23].

Materials and methods

Subjects

The data from 13 children (4 girls, 9 boys, mean age 11.0 years, range 7-13 years) with normal vision and without known neurological or psychiatric illness have been analyzed in the study. Previously we have used the data from this population to track the functional and morphological maturation of interhemispheric connectivity with a combined MRI technique [15]. Fourteen normal adults (6 women and 8 men; mean age 29.8 years, range of 21–42 years) without known neurological or psychiatric conditions and with normal or corrected-to-normal vision also participated in the study. The study followed the protocol approved by the Lausanne University Ethics Committee and written informed consent was obtained from children and their parents as well as from the adult participants. All the instrumental procedures conformed to the Declaration of Helsinki (1964) of the World Medical Association concerning experimentation on humans.

Stimuli

Subjects viewed bilateral sinusoidal black-and-white gratings with a spatial frequency of 0.5 Hz, a contrast of 70 %,



a size of a lateral patch of $11^{\circ} \times 19^{\circ}$, and drifting with a temporal frequency of 2 Hz. The stimuli either obeyed Gestalt grouping rules [iso-oriented colinear gratings (CG)] or violated them [non-colinear orthogonally oriented (NG)]. The stimulus conditions were alternated with background (uniform gray screen of the same space-averaged luminance as the stimuli) in a balanced pseudo-randomized order, five (for children) or six times (for adults), 15 s each (for a detailed description of the stimuli and eye movement control see [10, 11]).

fMRI data acquisition and pre-processing

For children, the protocol was performed on a 1.5 Tesla Siemens Magnetom Vision. Functional MRI images were acquired with an EPI gradient echo sequence (FA 90°, TE 66 ms, pixel size 3.75×3.75 mm², acquisition time 1.7 s, 22 slices, 5 mm thick) with a TR 3 s.

The adults were imaged on a 3T Siemens Trio scanner with the same type of sequence (FA 90°, TE 30 ms, pixel size 3×3 mm², acquisition time 1.7 s, 32 slices, 3 mm thick, TR 2 s). As the structural basis for brain segmentation and surface reconstruction, we acquired high-resolution T1-weighted 3D gradient echo sequence (MPRAGE), 160 slices ($1 \times 1 \times 1$ mm voxel size). In both experiments, we prevented head movements by cushioning the participant's head in a padded coil.

Both fMRI datasets were pre-processed and analyzed using Statistical Parametric Mapping (SPM8, Wellcome Department of Cognitive Neurology, London, UK), first, for standard single-subject and group statistics, then for DCM-specific design. Functional images were realigned to the first scan using a six-parameter rigid-body transformation and co-registered to the respective anatomical acquisition. To minimize inter-subject anatomical differences and increase the signal-to-noise ratio, each subject's images were normalized to the Montreal Neurological Institute template (MNI) using a 12-parameter affine transformation [24], re-sampled to $2 \times 2 \times 2 \text{ mm}^3$, and smoothed with an isotropic Gaussian kernel (FWHM = 6 mm).

It should be mentioned that moderately stronger and more extensive activations that can be recorded with the 3T compared to 1.5T scanner in a visual perceptual task [25] do not compromise the validity of comparative DCM analysis, since the method models the temporal dynamics of BOLD signals rather than their intensity. Furthermore, the local character of the modeled network situated within the visual cortex suggests the homogeneity of magnetic field across the regions of interest (ROIs). Finally, our ROIs (for details see "ROIs for DCM") are similarly located in the adult data from 1.5T [10, 11] and 3T (here), and, therefore, represent the same network.

Single-participant analysis was performed by adjusting the general linear model to our block design experiment. The signal drift across acquisitions was removed with a high-pass filter. Statistical parametric maps of the contrasts of interest were computed for each subject and were used as input values for the group statistics based on random field theory. In particular, the inferential statistics included between-condition t tests. To identify the nodes for DCM analysis (hereafter referred to as ROIs), we considered centers of clusters surviving a statistical threshold for peak height at P < 0.05 [family wise error (FWE) corrected] and satisfying an extent threshold of t > 30 contiguous voxels in the group SPMs.

Dynamic causal modeling

DCM serves to assess condition-dependent interactions within the distributed brain network based on an fMRI time series. To this end, DCM considers the brain as a dynamic system, in which external inputs cause changes in neuronal activity, which, in turn, cause changes in the distributed fMRI signals [9]. Mathematically, DCM for fMRI includes a bilinear model of neural dynamics and an extended balloon hemodynamic model of the transformation of neural activity into a measured BOLD response [9, 26]. A fixed set of ROIs, possible connections between them, and driving inputs are prerequisites for determining effective connectivity, which is defined by the influences that one node in a model exerts over another. Effective connections are divided into the intrinsic connections, i.e., the influences between ROIs in the absence of a modulating context, and the modulatory connections, i.e., the changes due to the experimental context. Effective connections can be positive, i.e., excitation in a source region induces increase in activity in the target, and negative, i.e., functional activation in the source causes target inhibition. Since the theoretical and practical issues of DCM were recently reviewed, we refer interested readers to these sources for details of the method and underlying theory [27–29].

Roles for DCM

The results of standard GLM analysis of SPMs reported earlier [10, 15] served as the basis for ROI selection. Both children and adults showed extensive activations in the striate and extrastriate areas under CG and NG conditions. In the primary visual cortex, we have chosen ROIs centered on the local group maxima from the conjunction maps of CG > background and NG > background contrasts. In the extrastriate areas, we defined ROIs from the CG > NG contrast, which revealed bilateral activations in the lingual gyrus (V3v, Fig. 1) in both groups. At the group level, the local maxima within these activated clusters served as ROI



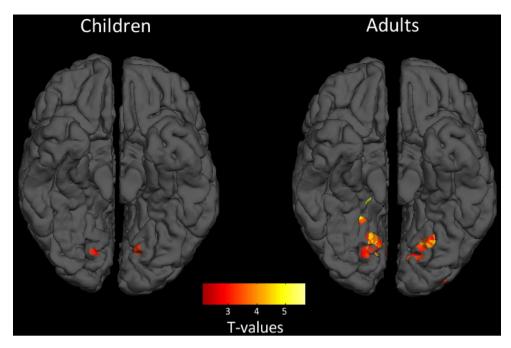


Fig. 1 Interhemispheric integration effects as revealed by fMRI activation. The 3D representation of statistical maps for the contrast between colinear and non-colinear gratings in the groups of children and adults (P < 0.001, extent threshold k = 30 voxels) is rendered on the MNI canonical brain. The *bottom view* of a pial surface is presented. In adults, the MNI coordinates (x, y, z) of the center of lefthemisphere cluster (178 voxels) are -12, -70, -10 and those of

Table 1 Anatomical locations of regions activated by gratings

Group	CG > background		CG > NG	
	Hemisphere			
	Left	Right	Left	Right
Adults	59 % V1, 41 % V2	66 % V1, 34 %V2	58 % V3v, 42 % V4	65 % V3v, 35 % V4
Children	62 % V1, 38 % V2	69 % V1, 31 % V2	51 % V3v, 49 % V4	70 % V3v, 30 % V4

Functional locations of the centers of activated clusters are given according to a probabilistic atlas [61]. The percentage represents each center's mean probability of belonging to the mentioned visual areas across subjects

centers. In individual subjects, the location of an ROI center was determined as a local maximum closest to the respective group location.

For each subject and ROI, the first principal component of the original time series of all voxels within a 4-mm radius sphere was extracted. In the primary visual cortex, ROIs were located at the V1/V2 border (Table 1). The ROIs in the extrastriate cortex occupied territory in V3v/V4. Hereafter, for the sake of simplicity, these two pairs of ROIs are referred to as V1_L, V1_R, V3_L, and V3_R, where the "L" and "R" designate the left and right hemispheres, respectively.

right-hemisphere cluster (133 voxels) are 16, -76, -14. In children, the coordinates are -18, -78, -14 (left hemisphere, 88 voxels) and 10, -76, -6 (right hemisphere, 95 voxels). Hot scale shows T values. In both groups, BOLD response is located within the lingual gyri, but the clusters are larger and more significant in the adult group. The centers of the clusters (P < 0.05, FWE corrected) define the V3 locations for DCM analysis

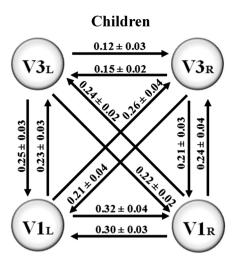
Dynamic causal model

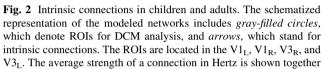
For the sake of compatibility between children and adults, we applied a single model to both groups (Fig. 2). This is a fully connected model of intrinsic connections, which has also been a winning one for the group of aged healthy adults under the same perceptual task [22]. In this model, the stimuli > background contrast serves as a driving input. The signals induced by the bilateral stimuli travel via crossed visual pathways and directly affect the left and right primary visual cortices. The induced activity then spreads through the system according to the intra- and interhemispheric reciprocal connections. On the assumption that each intrinsic connection can be modulated by the task, we reproduced the architecture of intrinsic connections for modulatory connections (Fig. 3). We modeled the CG > stimuli contrast, since in our experiment the interhemispheric integration is a source of modulatory effects.

Group statistics for DCM

To test for differences in the coupling parameters between developing and mature brain networks, we applied mixed between-within analysis of variance (ANOVA). The developmental affects were analyzed with a between-







with a standard error next to the respective connection. For the sake of simplicity, the driving inputs provided by the right visual field gratings to Vl_L and by the left visual field gratings to Vl_R are not shown

 $m V3_{R}$

Adults

 0.11 ± 0.02

 0.10 ± 0.02

 0.21 ± 0.03

subject age factor (children vs. adults). According to our hypothesis, developmental effects might differ depending on the topography of connections, which, in terms of ANOVA, can be shown with an interaction of the topography of connections with the age factor. The planned between-group comparisons of connection parameters were performed by means of a two-sample two-tailed t test. For descriptive statistics of each parameter, we used a one-sample two-tailed t test. Significant effects are reported at P < 0.05 corrected for multiple comparisons with the Bonferroni method.

The links between the coupling parameters of the DC model and morphometric measurements of the corpus callosum (CC) were assessed by the Pearson correlation coefficient ρ with corresponding P value. All the statistical tests were implemented with SPSS 17.0 for Windows.

Results

Intrinsic connections and driving inputs do not change with age

Group-averaged estimates of the strength of intrinsic connections are shown in Fig. 2. All the parameters significantly differ from zero at P < 0.05 (two-tailed t test, Bonferroni corrected). To explore the age-related changes in vertical effective connections between striate (V1) and extrastriate (V3) areas, including both intra- and interhemispheric connections, we applied a three-way between-within ANOVA with factors of age (children vs. adults), direction (top-down vs. bottom-up), and topography (4)

levels: $V1_L - V3_L$, $V1_L - V3_R$, $V1_R - V3_L$, and $V1_R - V3_R$). Neither the main effect of age, nor its interactions with the other two factors proved significant. We also failed to find age-related changes in horizontal interhemispheric connections at the striate and extrastriate levels, within the framework of a three-way betweenwithin ANOVA with factors of age (children vs. adults), hierarchy (V1 vs. V3), and direction $[V1(3)_L \rightarrow V1(3)_R]$ vs. $V1(3)_L \leftarrow V1(3)_R]$.

Similarly, we found no differences in driving inputs between the groups or hemispheres as shown with a twoway between-within ANOVA with factors of age (children vs. adults) and hemisphere (left vs. right).

Modulatory connections strengthen with age

All the modulatory connections turned out to be inhibitory (Fig. 3). Six out of 12 connections in children and all but one in adults significantly differ from zero at P < 0.05(two-tailed t test, Bonferroni corrected). Note that significant connections in children are mostly bottom-up. The vertical modulatory connections were analyzed with a twoway between-within ANOVA, which includes the age and topography factors described for intrinsic connections. The main effect of age was significant at P = 0.037 (F = 4.88, df = 1, $\eta^2 = 0.163$). As Fig. 3b shows, the strength of inhibitory top-down connections builds up with age. A three-way between-within ANOVA with the age, hierarchy, and direction factors (described for horizontal intrinsic connections) revealed the main effect of age (P < 0.001,F = 18.289, df = 1, $\eta^2 = 0.422$) for horizontal modulatory connections, which appeared to be stronger in adults.



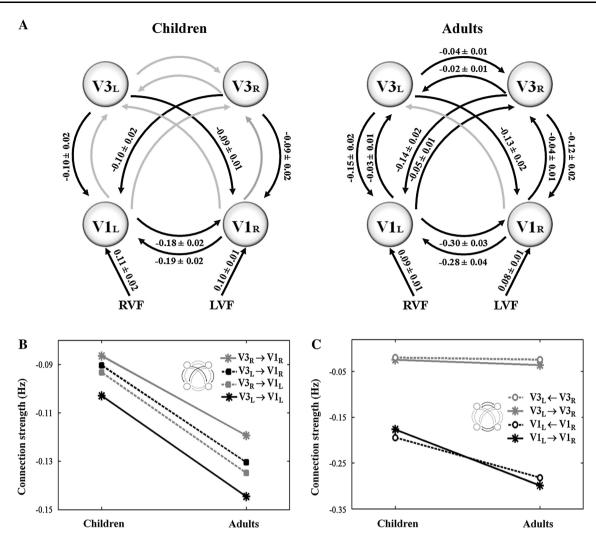


Fig. 3 Modulatory connections for children and adult groups. a The average estimates of modulatory parameters in Hertz with standard errors are shown alongside the respective connections. Other designations are as in Fig. 2. b The main effect of age on top-down

modulatory connections. Connections from left and right hemispheres are in *black* and *gray*, respectively. **c** The main effects of the age factor and the interaction of age \times hierarchy on the striate (in *black*) and extrastriate (in *gray*) interhemispheric connections

This effect interacted with a hierarchy factor at P=0.007 (F=8.589, df=1, Greenhouse–Geisser corrected; $\eta^2=0.256$). The planned between-group comparisons showed that only striate connections are significantly stronger in adults than in children (P=0.001, $V1_L \rightarrow V1_R$; P=0.013, $V1_L \leftarrow V1_R$, Fig. 3c).

Effective coupling is associated with brain maturation

To visualize the developmental trajectory for interhemispheric effective connectivity and to link it to structural brain maturation, we applied a polynomial regression analysis to the effective connections between primary visual areas that were significantly stronger in adults than in children. To this end, age and the CC area assessed on the basis of a midsagittal slice served as estimators of general development and of structural maturation,

respectively. The average strength of the $V1_L \rightarrow V1_R$ and $V1_L \leftarrow V1_R$ modulatory connections was used as a dependent variable. We found that both linear and quadratic regression models are significant for age (P=0.002; Fig. 4a) and for the CC area (P=0.004 for the linear and P=0.001 for the quadratic model; Fig. 4b). The quadratic models showed better fit to the data (41.1 and 44.9 % of variance for age and CC area, respectively) than the linear models (34.3 and 30.3 %).

Although the CC area correlates with age ($\rho = -0.62$, P = 0.001), the latter is a more general index, accumulating all the developmental changes. To assess the contribution of the individual variations of the CC area, while controlling for age, we applied hierarchical multiple regression analysis. Preliminary analysis ensured no violations of the assumptions of normality, linearity, and multi-colinearity (tolerance = 0.62). In the hierarchical



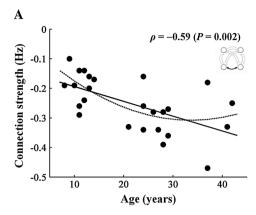


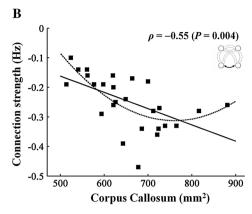
Fig. 4 Effective connectivity and structural brain maturation. The *plots* show scatter diagrams and developmental trajectories fitted with the first- and second-level polynoms (*solid* and *dashed line*, respectively) for age (**a**) and the CC area (**b**). Pearson regression coefficient ρ with corresponding P value represents a measure of bivariate

regression model with the two variables, which turned out to be significant at P = 0.003 and explained 40 % of the variance in connection strength, the CC area predicted only 5.7 % in addition to what age predicted. Therefore, being controlled for the effect of age, the CC area failed to explain a significant amount of variance.

Discussion

Development of spatial integration and changes in top-down effective connections

In children, the key mechanisms of spatio-temporal integration are functional beginning in infancy. In particular, newborns looking at point-light animations prefer biological motion to non-biological motion and, among the biomotion animations, distinguish between upright and inverted displays [30]. Beginning at 2 months of age, infants combine the coherently moving components across large areas of space and treat them as connected wholes [31]. By 3–4 months of age, they are able to perceive the global structure of various arrays [32]. Further development of visual spatial integration continues through middle childhood and, probably, adolescence. Sensitivity to the global structure of Glass patterns with a different percentage of noise approximates that of adults at 9 years of age [33]. The capacity to integrate spatially distant line segments based on their colinearity improves until 10 years of age [34]. Spatial integration tested in contour-detection tasks improves until about 14 years of age [12]. A similar trajectory was shown for sensitivity to biological and global motion [35]. Furthermore, the processing of global shape in paradigms including compound displays still differs in adolescence from that in adults [36, 37].



correlation. The CC area was assessed from a midsagittal slice. The connection strength refers to the average value of reciprocal modulatory connections between the primary visual (*black line* on the connection scheme)

These behavioral changes are paralleled by the development of functional connectivity in neural networks as manifested by EEG synchronization [38–41]. To show the transformations of neural networks underlying the protracted maturation of integrative mechanisms, here we have applied the DCM method to an fMRI time series under a passive perceptual task, which contrasted bilateral colinear gratings (Gestalt) to bilateral orthogonal gratings (no Gestalt), of which only the first stimulus can be interhemispherically integrated.

Our analysis is implemented in the framework of predictive coding theory, which provides an elegant model of the hierarchical optimization mechanism. It is based on feedback connections that carry predictions of lower-level neural activity and feed-forward connections that carry residual errors between predictions and actual lower-level responses [8, 17]. Indeed, a DCM analysis of fMRI activations induced by coherent or meaningful stimuli compared to random ones has shown inhibitory modulatory effects from the higher-order to the lower-order visual areas in adult subjects [21].

Here, we have shown that intrinsic effective connections within the visual areas do not differ between children older than 7 years and young adults, suggesting that basic visual networks integrated via long-distance reciprocal excitatory pathways are established at this age. The intrinsic connections were modulated by the coherent CG stimulus both in children and in adults. All the descending and lateral modulatory connections turned out to be inhibitory. Therefore, here we have shown for the first time that inhibitory backward effective coupling recently reported in adults [21, 22] works in children as well.

This result agrees with a large body of literature reporting the higher-order effects on V1 via feedback connections originating within the visual system and/or via



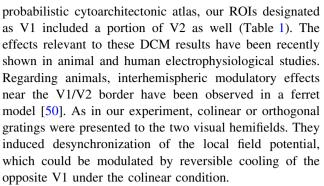
long-range local horizontal connections [42, 43]. Although low-level integration has been predominantly studied intrahemispherically as a contextual modulation in the neighborhood of the classical receptive field [44, 45], in adults it likely operates at a long range as well, e.g., across the cerebral hemispheres. In this regard, Ban and colleagues have demonstrated that interhemispheric integration can result in changes of V1 activation [20]. They have found a significantly lower BOLD response to two arcs located symmetrically in the lower visual field quadrants (i.e., to a pattern following the Gestalt principle of good continuation) compared to their asymmetrical (diagonal) location. In the absence of direct interhemispheric V1 connections between the low and high visual quadrants, this modulation of V1 activation is likely due to the topdown influences from the extrastriate areas.

The finding that the top-down modulatory effects in adolescents differ from those in adults suggests their slow formation in human ontogenesis. The available data on the structural maturation of feedback connections, which might be a part of the neural network that enables colinearity detection in humans [12], are limited to those between V2 and V1 [1, 2]. According to this postmortem anatomical study, the upper layers of V1, which receive the feedback connections, seem to be immature even at 5 years of age. In contrast, we found no evidence supporting the immaturity of effective connections downstream of V1, which have also been proposed to underlie the slow progression of perceptual integration in human ontogenesis [35].

The abnormal development of top-down connections in children can explain a well-known interictal phenomenon of reduced habituation to repetitive visual and auditory stimuli in migraine patients [46–48]. Sensory habituation, characteristic for healthy adults and children, refers to progressive reduction in the cortical response to repetitive stimuli. It is manifested as a gradual decrease in the amplitude of evoked potentials. Habituation can be considered as a mechanism protecting against the accumulation of potentially toxic metabolites in the brain. According to the predictive coding hypothesis, the repetition reduces cortical response due to increasing predictability of successive stimuli emerging as a direct result of predictive processing in the brain [49]. Since this effect is sustained through inhibitory feedback connections, their formation in children, which suffer from migraine, should be thoroughly explored in future studies.

Interhemispheric modulation in V1

In addition to feedback connections strengthening with age, we found increasing interhemispheric inhibitory effects in the primary visual cortex. More precisely, according to the



In adult humans, interhemispheric inhibition in the primary visual cortex has been shown with unilateral transcranial magnetic stimulation of V1, which increased (by means of disinhibition) the amplitude of EEG potentials, evoked by gratings, in the contralateral hemisphere [51]. Another example of transcallosal inhibition in the human primary visual cortex has been demonstrated in binocular rivalry. During perceptual transitions, a dominant stimulus can spread across the visual field in a wave-like manner. These traveling waves of rivalry dominance are delayed when passing from one visual hemifield to another. In a diffusion tensor imaging study, the diffusion properties of callosal fibers connecting the left and right V1 reliably predicted this across-hemifield delay [52]. Since perceptual alternation during binocular rivalry is likely to result from inhibitory interactions between neural representations of the different percepts [53], this finding links the CC to interhemispheric inhibition.

Hypothetically, the insufficiency of interhemispheric inhibition between the primary visual areas could be among the factors behind the reduced habituation (the increased amplitude of visual evoked potential) in patients with migraine. However, until now only the late components of visual evoked potentials, generated in the extrastriate areas, are shown to be different (e.g., [54]).

From an ontogenetic perspective, immature transcallosal modulation between the primary visual areas in the absence of age-dependent interhemispheric effects in the extrastriate areas challenges the conventional view that posits the prior maturation of the early visual cortex as a precondition for the later development of higher-order ventral stream regions [55]. Yet our finding is supported by the association between the indices of effective connectivity and brain maturation. The interhemispheric interaction in the ventral visual stream is implemented via fibers that form the splenium of the CC [56, 57]. Poorly myelinated fibers of small diameter with a protracted course of myelination are abundant in this region [58]. The CC area depends on the number of axons, their size, and the thickness of the myelin sheaths [15, 59], providing a reasonable cumulative index of interhemispheric connectivity. As we found, its agedependent changes explain about 40 % of the variance in



interhemispheric effective inhibition between the primary visual areas.

Since interhemispheric effective inhibition is presumably implemented via polysynaptic pathways with long-distance excitatory and local inhibitory components, this correlation reflects only the development of the long-distance part. As far as local inhibitory mechanisms in the primary visual cortex are concerned, they were recently analyzed postmortem [60]. Consistently with our results, this study revealed the protracted development of GAB-Aergic mechanisms, which continues well into the second and third decades of life.

In sum, our results suggest that feedback connections from the extrastriate to the primary visual areas together with the interhemispheric connections between primary visual areas strongly contribute to spatio-temporal feature integration in the mature brain. Both feedback and horizontal effective connectivity have a long developmental trajectory extending into the adolescence. This wide developmental window suggests a protracted period of the vulnerability to neuropathological disorders including migraine. For better understanding of the mechanism inducing migraine, the DCM can be used. The method seems to be especially efficient for the analysis of brain networks providing habituation.

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