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1 **Data-Driven Classification of Spectral Profiles Reveals**

2 **Brain Region-Specific Plasticity in Blindness**

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6 Running title: Region-specific Spectral Plasticity

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1 **Abstract**

2 Congenital blindness has been shown to result in behavioral adaptation and neuronal reorgani-
3 zation, but the underlying neuronal mechanisms are largely unknown. Brain rhythms are char-
4 acteristic for anatomically defined brain regions and provide a putative mechanistic link to cog-
5 nitive processes. In a novel approach, using magnetoencephalography resting state data of con-
6 genitally blind and sighted humans, deprivation-related changes in spectral profiles were
7 mapped to the cortex using clustering and classification procedures. Altered spectral profiles in
8 visual areas suggest changes in visual alpha-gamma band inhibitory-excitatory circuits. Re-
9 markably, spectral profiles were also altered in auditory and right frontal areas showing in-
10 creased power in theta-to-beta frequency bands in blind compared to sighted individuals, pos-
11 sibly related to adaptive auditory and higher-cognitive processing. Moreover, occipital alpha
12 correlated with microstructural white matter properties extending bilaterally across posterior
13 parts of the brain. We provide evidence that visual deprivation selectively modulates spectral
14 profiles, possibly reflecting structural and functional adaptation.

15

16 ***Keywords:*** congenital blindness; MEG; oscillations; spectral fingerprints

17

18

1 **Introduction**

2 Brain rhythms occur ubiquitously across cortex (Buzsáki 2004; Buzsáki et al. 2013; Singer
3 2018) and have been related to cognitive functions. Investigating changes in brain rhythms in
4 congenital blindness can provide crucial insights into the neuronal mechanisms of behavioral
5 changes as found in this population. Congenitally blind individuals perform better on various
6 auditory, tactile, and higher-cognitive tasks compared to sighted controls (Bull et al. 1983; Les-
7 sard et al. 1998; Roeder et al. 2001; Amedi et al. 2003; Gougoux et al. 2005; Foecker et al.
8 2012), including tasks where certain aspects of temporal processing are essential, such as in
9 temporal order processing tasks, task involving musical meter or ultra-fast speech processing
10 (Hoetting et al. 2004; Roeder et al. 2004; Stevens and Weaver 2005; Moos and Trouvain 2007;
11 Roeder et al. 2007; Trouvain 2007; Hertrich et al. 2009; Dietrich et al. 2013; Lerens et al. 2014;
12 Carrara-Augustenberg and Schultz 2019; Zhang et al. 2019). Improved behavioral performance
13 in auditory or tactile tasks in congenitally blind individuals is accompanied by cross-modal
14 plasticity. That is, the “visual” cortex has been observed to be recruited during non-visual tasks
15 (Burton 2003; Pascual-Leone et al. 2005; Bedny et al. 2011; Voss and Zatorre 2012; Gudi-
16 Mindermann et al. 2018; Rimmele et al. 2019), which has been postulated to functionally con-
17 tribute to compensatory task performance. Additionally, intramodal plasticity has been ob-
18 served in congenitally blind individuals. That is, the auditory (Elbert et al. 2002; Stevens and
19 Weaver 2005; Gougoux et al. 2009; Schepers et al. 2012; Hertrich et al. 2013; Watkins et al.
20 2013) and somatosensory (Pascual-Leone and Torres 1993; Roeder et al. 1996; Sterr et al.
21 1998a; Sterr et al. 1998b) cortices were reported to show altered (mostly increased) neuronal
22 activity during task processing. Intramodal plasticity has been related to early postnatal sensory
23 experiences shaping the functional development of cortical representations as shown in animal
24 research (Zhang et al. 2001; for review: Rauschecker 2008). For instance, reorganization of
25 primary somatosensory and auditory areas has been reported in cats that were visually deprived

1 from birth (Rauschecker et al. 1992; Korte and Rauschecker 1993). Although these behavioral
2 and neuronal adaptations have been observed, the mechanisms underlying visual deprivation-
3 related plasticity are still poorly understood.

4 Studying brain rhythms in congenitally blind individuals might provide further insights into the
5 mechanisms of neuronal plasticity, as brain rhythms reflect neuronal circuitry activity and have
6 been related to various sensory, motor and higher-cognitive processes, as discussed in the fol-
7 lowing. Brain rhythms are thought to reflect the synchronization (phase-alignment) of oscilla-
8 tory activity across neuronal populations, subserving the formation of both local assemblies and
9 large-scale functional networks (Engel et al. 1992; Hipp et al. 2011; Raichle 2011; Siegel et al.
10 2012; Singer 2013). Specific spectral profiles have been conjectured to be associated with dif-
11 ferent anatomical areas (Giraud et al. 2007; Keitel and Gross 2016). Such spectral profiles are
12 assumed to reflect the intrinsic brain rhythms of a brain area that are related to distinct processes
13 (Keitel and Gross 2016). Importantly, intrinsic brain rhythms measured spontaneously (e.g.,
14 ongoing activity during resting state) reflect the functional organization of the brain and thus
15 the default architecture recruited during task-related processes (Smith et al. 2009; Deco et al.
16 2011; Raichle 2011; Engel et al. 2013; Raichle 2015; Keitel and Gross 2016; Sormaz et al.
17 2018) because intrinsic brain rhythms oscillate in a characteristic way even when not involved
18 in task-related processing (Brookes et al. 2011; Deco et al. 2011). For example, Keitel and
19 Gross (2016) have shown that a characteristic pattern of delta-, theta-, alpha-, and beta-band
20 activity is present in primary auditory cortex both during rest and speech comprehension. While
21 the global appearance of the spectral profile of primary auditory cortex was similar during
22 speech comprehension and rest, the relative contribution of individual frequency clusters (i.e.,
23 their spectral power) was altered by the condition. During speech comprehension, in compari-
24 son to rest, the alpha band was suppressed, indicating reduced inhibition of sensory cortices

1 (Jensen and Mazaheri 2010). The other bands, which were associated with several levels of
2 speech processing, were enhanced.

3 Various studies have observed that brain rhythms are recruited in a task-specific manner during
4 sensory, motor and higher-cognitive tasks, linking intrinsic brain rhythms to cognitive functions
5 (Engel et al. 1992; Singer and Gray 1995; Engel et al. 2001; Lakatos et al. 2008; Holcombe
6 2009; Giraud and Poeppel 2012; Lakatos et al. 2013; Heusser et al. 2016; Portoles et al. 2018;
7 Rimmele, Gross, et al. 2018; Schroeder et al. 2018; VanRullen 2018). For instance, in sighted
8 individuals, the role of the alpha rhythm (~8–12Hz) in cognition is well-established, with pos-
9 terior alpha- and gamma-band oscillations coupling in excitatory-inhibitory cycles (Klimesch
10 et al. 2007; Buffalo et al. 2011; Jensen et al. 2012). Specifically, the alpha rhythm mediates the
11 inhibition of task-irrelevant neuronal circuits. Gamma-band oscillations (30–100 Hz) in occip-
12 ital cortex, by contrast, allow for feedforward processing of task-relevant visual information
13 (van Kerkoerle et al. 2014; Michalareas et al. 2016). In congenitally blind individuals, a reduced
14 or absent occipital alpha rhythm has repeatedly been observed (Adrian and Matthews 1934;
15 Noebels et al. 1978; Kriegseis et al. 2006; Hawellek et al. 2013; Schubert et al. 2015). The lack
16 of visual input during a sensitive period of brain development in congenitally blind individuals
17 presumably results in atrophy or reorganization of posterior alpha generators (Ptito et al. 2008;
18 Wang et al. 2013; Aguirre et al. 2016; Reislev et al. 2016). Posterior alpha generators have been
19 located within the cortical and thalamo-cortical visual pathways (Lopes da Silva 1991; Lőrincz
20 et al. 2009) and posterior gamma generators within the cortical visual pathways (Bastos et al.
21 2014; Marshall et al. 2018). Beyond the occipital brain rhythms, however, no clear hypothesis
22 linking brain rhythms and structural alterations in congenitally blind individuals can be made.

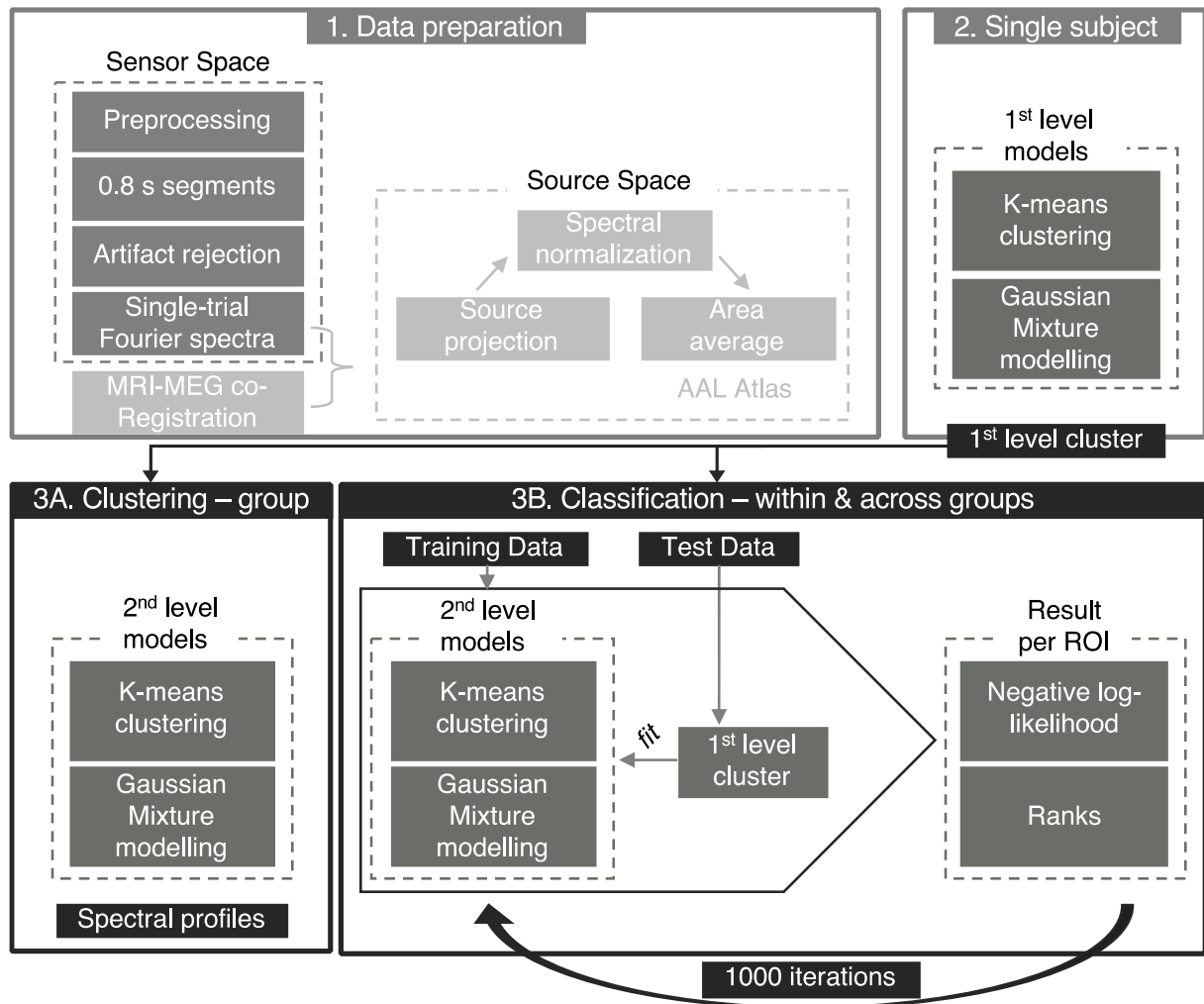
23 In sighted individuals, beyond the alpha rhythm, intrinsic theta-band oscillations (~4–7 Hz) in
24 auditory cortex (Lakatos et al. 2005; Keitel and Gross 2016) have been widely studied. Theta
25 brain rhythms likely play a central role in the temporal segmentation of speech signals during

1 speech comprehension (Luo and Poeppel 2007; Giraud and Poeppel 2012; Gross et al. 2013;
2 Doelling et al. 2014; Pittman-Polletta et al. 2020), as well as segmentation of music and envi-
3 ronmental sounds (Henry et al. 2014; Doelling and Poeppel 2015). Interestingly, initial findings
4 have linked the ability of congenitally blind individuals to comprehend ultra-fast speech—at
5 rates where speech comprehension fails in sighted individuals—to accelerated theta brain
6 rhythms in auditory cortex, as well as the recruitment of additional brain areas such as the visual
7 cortex (Trouvain 2007; Hertrich et al. 2009; Dietrich et al. 2013; Hertrich et al. 2013; see also:
8 Van Ackeren et al. 2018). Furthermore, initial research suggests that other brain rhythms in
9 brain areas outside of visual cortex might also be altered in congenitally blind compared to
10 sighted individuals (e.g., such as the beta- and gamma-band rhythm during working memory or
11 semantic categorization tasks, as indicated by functional connectivity measures: Schepers et al.
12 2012; Gudi-Mindermann et al. 2018; Rimmele et al. 2019). However, the mechanisms under-
13 lying these spectral alterations remain to be elucidated. In summary, the systematics of how
14 spectral profiles outside of visual cortex are altered due to congenital visual deprivation, are not
15 well understood and no clear hypothesis can be derived beyond the occipital alpha rhythm.

16 The goal of the present study is to characterize alterations in spectral profiles in congenitally
17 blind as compared to sighted individuals in order to map visual deprivation-related spectral
18 changes to brain areas. By studying a broad range of frequencies, instead of narrow frequency
19 bands, our research provides the means towards linking changes in spectral profiles across the
20 whole brain to adaptive behavior in congenitally blind individuals in future research. Standard
21 analyses of brain rhythms face several problems (see Keitel and Gross 2016), which possibly
22 explains the lack of a comprehensive account of spectral changes in congenitally blind individ-
23 uals beyond the alpha rhythm. Predominant neuronal activity in the alpha and super-low fre-
24 quency ranges ($1/f$) complicates the analysis of spectral properties in the lower frequency
25 ranges. Furthermore, the brains' temporal dynamics are often captured poorly over the course

1 of the recording session (de Pasquale et al. 2010; Singer 2013). Here, we have employed a novel
2 analysis pipeline, introduced by Keitel and Gross (2016), which reveals brain area-specific
3 spectral profiles. The analysis pipeline overcomes the limitations of standard analyses of brain
4 rhythms by using segment-based clustering (of source-localized Fourier spectra) to disentangle
5 spectral properties in the lower frequency ranges and by taking the time course of activity into
6 account. We extended the pipeline (Fig. 1) to quantify differences in spectral profiles across
7 cortical brain areas between congenitally blind and sighted individuals by using a cross-classi-
8 fication approach. This approach allowed us to identify brain areas with changed spectral pro-
9 files as a result of congenital blindness. More specifically, the clustering procedure reveals
10 multi-dimensional spectral profiles that might consist of several clusters reflecting distinct in-
11 trinsic brain rhythms.

12 First, we expected to replicate Keitel and Gross (2016) by showing that spectral profiles in
13 sighted adults are region-specific, enabling the classification of brain regions based on spectral
14 profiles. Second, we hypothesized that within a group of congenitally blind individuals, spectral
15 profiles would similarly follow specific patterns and enable the classification of brain regions.
16 Third, visual deprivation-related plasticity was predicted to result in altered spectral profiles in
17 congenitally blind individuals as compared to the sighted, particularly for brain regions for
18 which visual deprivation-related reorganization (cross- and intra-modal plasticity) has previ-
19 ously been shown, such as for sensory cortices. A specific hypothesis regarding the changes in
20 brain rhythms in congenitally blind individuals was only derived for the alpha rhythm, which
21 was expected to be reduced in occipital cortex. Finally, we expected a relationship of micro-
22 structural white matter properties and spectral profiles of occipital cortex.



1

2 **Figure 1. Analysis pipeline adapted from Keitel and Gross (2016).** (1) Continuous resting state MEG data were
 3 preprocessed and segmented into trials of 0.8 s length. Complex Fourier spectra were computed for each trial
 4 separately and projected into source space using previously defined beamforming (LCVM) coefficients. The data
 5 were spatially normalized, dividing each voxel's power by the mean power of all trials and voxels. Voxels were
 6 grouped according to the AAL atlas and power values were averaged across voxels of each anatomical area ($N =$
 7 115). (2) In the 1st-level analysis, power matrices were clustered into 9 distinct spectral clusters per participant and
 8 brain region using k-means and Gaussian Mixture model algorithms. (3A) In the 2nd-level analysis, 1st-level indi-
 9 vidual clusters were again subjected to k-means clustering and GMMs to establish region-specific spectral clusters
 10 consistent at the group level, also referred to as spectral fingerprints. The optimal number of group-level clusters
 11 per anatomical area was defined by the Silhouette Criterion prior to the group-level clustering procedure. (3B) For
 12 the classification procedure the experimental group was divided into training and test set. For each brain region,
 13 the fit between 1st-level clusters of the test group and group-level clusters of all regions of the training set was
 14 calculated. For each anatomical region, this resulted in a negative log-likelihood value for all regions (i.e., 115
 15 values per region), indicating its similarity to all brain regions based on the spectral clusters. This fitting procedure
 16 between training and test set was repeated 1000 times with new group assignment (training vs. test) on each iter-
 17 ation. The interindividual variance within test and training groups of each iteration was controlled for with an ad-
 18 ditional 100 iterations within the respective sets.

1 **Materials and Methods**

2 **Participants**

3 The study was approved by the German Psychological Association. All participants gave writ-
4 ten informed consent prior to the experiments and received monetary compensation. The data
5 were recorded in the context of a larger project (Gudi-Mindermann et al. 2018; Rimmele et al.
6 2019). Three to four minutes of resting state MEG data were collected from a group of sighted
7 and congenitally blind individuals matched in age, gender and education. During data collection
8 the congenitally blind (CB) and the sighted (S-BF) participants were blindfolded. The blind-
9 folding of CB participants aimed at reducing involuntary eye movements. Additionally, for the
10 sighted a resting state measurement with open eyes was conducted (S-EO). The data reported
11 here include (after a few subjects were excluded, see below) 26 subjects for the CB (12 females;
12 mean age: 37.8 years; *SD*: 10.2 years; age range: 22–55 years), 24 for the S-BF (11 females;
13 mean age: 36.8 years; *SD*: 10.1 years; age range: 21–55 years) and 23 for the S-EO (11 females;
14 mean age: 37.3 years; *SD*: 9.8 years; age range: 21–55 years). A few subjects were excluded
15 after data collection because of corrupted resting state files (one subject for the CB, one subject
16 for the S-EO) or lack of individual structural MRI scan (three subjects for the S-BF and S-EO).
17 All participants were healthy with normal hearing (self-report) and assured no history of psy-
18 chiatric or neurological disorders. One blind participant reported a history of a depressive mood
19 disorder, but was free of symptoms and without current treatment. Sighted participants had
20 normal or corrected-to-normal vision (self-report). In the blind, vision loss was total and re-
21 sulted from a variety of peripheral (pre)natal conditions (retinopathy of prematurity: $n=9$; ge-
22 netic defect, $n=5$; congenital optic atrophy: $n=2$; Leber's congenital amaurosis: $n=2$; congenital
23 cataracts, glaucoma: $n=2$; congenital retinitis: $n=2$; binocular anophthalmia: $n=2$; retinitis pig-
24 mentosa: $n=1$; congenital degeneration of the retina, $n=1$). Seventeen participants reported min-
25 imal residual light perception.

1 **MRI and MEG Data Acquisition**

2 For all participants T1-weighted structural MRI scans and DWI-MRI scans were obtained with
3 a 3T scanner (Siemens Magnetom Trio, Siemens, Erlangen, Germany). For the T1-weighted
4 images we used the following parameters: TE = 2.98 ms, TR = 2300 ms, flip angle = 9, and
5 isotropic 1 mm³ voxels, 256 sagittal slices. The MEG data were recorded in a magnetically
6 shielded room using a 275-channel whole-head system (Omega, 2000, CTF Systems Inc.),
7 while participants sat in an upright position. The data were acquired with a sampling rate of
8 1200 Hz. Prior to each experiment, the head position was measured relative to the MEG sensors
9 and during the recording the head position was tracked.

10 **Data Analysis**

11 The initial analyses in this study are adopted from the analysis pipeline proposed by Keitel and
12 Gross (2016). The modifications to the analysis pipeline and the novel analyses are stated in
13 detail below. All analyses were carried out using Matlab R2018a (version 9.4.0.813654, The
14 Math Works, Inc.), the Fieldtrip Toolbox (version 20181104), and SPM12.

15 **Data Preparation in Sensor Space: Preprocessing, Artifact Rejection, Source Localization**

16 During preprocessing, the MEG signal was downsampled to 250 Hz, denoised and detrended.
17 To better capture the dynamically changing spectral properties of the brain, the continuous sig-
18 nal was segmented into trials of 0.8 s. Trials were declared as noisy and excluded when their z-
19 score was higher than 2. On average, seven trials were excluded, resulting in a mean of 340.3
20 trials ($SD= 34.7$) per subject (S-EO: mean = 346.7, $SD = 37.4$; S-BF: mean = 336.8, $SD = 34.2$;
21 CB: mean = 338, $SD 33.2$). Due to shorter recordings in the present study, trial duration was
22 slightly shorter, relative to the 1 s duration used in Keitel and Gross (2016), to increase the
23 amount of trials. MEG channels were labeled as noisy and rejected when the ratio between their
24 noise level (in SD) and that of the neighboring sensors (in SD) exceeded a value of 0.5 ([Sensor

1 $SD - \text{Neighbor } SD] / \text{Neighbor } SD$; mean number of excluded channels = 1.22, $SD = 1.34$).
2 Finally, using independent component analysis (ICA), data was cleaned from heartbeat, eye
3 blinks and eye movements-related artifacts (components were identified based on their time-
4 course, topography and variance across trials). To prepare the source projection of the Fourier
5 spectra, beamformer coefficients were obtained. For this purpose, we applied co-registration of
6 individual T1-weighted MRI scans and the MEG coordinate system, realignment, segmentation
7 and normalization to Montreal Neurological Institute (MNI) space. A forward model was cre-
8 ated using a single-shell model and linearly constrained minimum variance (LCMV) beam-
9 former coefficients (Van Veen et al. 1997) were calculated for the MEG time series for each
10 individual voxel on the 10 mm regular grid.

11 **Spectral Analysis in Sensor Space**

12 The analyses described in the following were performed for all three groups separately (CB, S-
13 EO, S-BF; for an overview, see Fig. 1). First, Fourier spectra were calculated on 0.8 s long trials
14 for each subject, using a multitaper approach (3 tapers) and zero-padding (length of 2 s). We
15 slightly reduced the duration of the temporal segments compared to the duration used by Keitel
16 and Gross (2016), i.e., 1 s, in order to match the overall number of segments used for the anal-
17 yses while accounting for the shorter overall MEG recording time in our study. Second, using
18 the previously computed LCMV coefficients, the complex Fourier spectra were projected into
19 source space. Neuronal power, i.e., the squared amplitude of the Fourier coefficient, was com-
20 puted for each time segment, voxel, and frequency. Furthermore, neuronal power of individual
21 voxels and segments was ratio normalized, i.e., divided by the mean power across all voxels
22 and trials. This ratio normalization resulted in voxel-specific spectral properties with values
23 above/below 1 highlighting the differences of a given voxel to the mean spectral power across
24 all voxels separately at each frequency. All values were subtracted by 1 (leading to values
25 above/below zero), to facilitate the identification of changes in power (decreases/increases). We

1 additionally conducted a control analysis without the normalization by performing the cluster-
2 ing and classification procedures on the non-normalized data.

3 **k-Means Clustering and Gaussian Mixture Modelling of Source-Localized Spectral Ac-** 4 **tivity**

5 To identify region-specific spectral clusters in the individual subject, the brain was parcellated
6 according to the AAL atlas (Tzourio-Mazoyer et al. 2002; 116 regions of interest, ROIs). For
7 one anatomical region (cerebellum 3L), however, the interpolation between the AAL atlas and
8 the source model was not successful. Thus, this region was excluded and all analyses are based
9 on the remaining 115 anatomical areas. For each of the ROIs, voxels were grouped and power
10 spectra were averaged across voxels. Within a brain area, neuronal populations might express
11 several distinct intrinsic brain rhythms, which together constitute the spectral profile of an area.
12 Clustering algorithms were employed to identify spectral clusters, reflecting intrinsic brain
13 rhythms. First, trial-by-frequency matrices were subjected to a k-means algorithm (MacQueen
14 1967) which established spectral clusters by partitioning the n observations (0.8 s temporal
15 segments) into k clusters. For the 1st-level analysis, the k was set to 9, based on the Silhouette
16 criterion evaluation (Rousseeuw 1987). Second, for each subject and ROI, Gaussian Mixture
17 Models (GMMs; Reynolds and Rose 1995) were fitted to the 9 clusters obtained from the k-
18 means analysis (1st-level GMM). Next, in order to identify the optimal number of clusters per
19 brain region across all subjects for the 2nd-level group analysis, the 1st-level GMMs were eval-
20 uated using the Silhouette criterion. Silhouette values were computed for cluster solutions in
21 the range from 1 to 15, the fitting was repeated 1000 times. At the group level, k-means clus-
22 tering was applied to the 1st-level clusters in order to disclose consistent patterns across subjects.
23 The optimal number of clusters per brain area, as assessed by the Silhouette criterion evaluation
24 (Rousseeuw 1987), was used as k-parameter for the algorithm. As before, k-means results were
25 fed into GMM revealing the final clusters per brain region (2nd-level GMM).

1 Clusters were considered for visualization only if they were reflective of the majority of partic-
2 ipants. To facilitate reading of the spectral plots, group-level clusters were color-coded accord-
3 ing to the frequency of the maximum amplitude of the cluster (peak frequency) (delta: 1–3.5
4 Hz, red; theta: 4–8 Hz, green; alpha: 8.5–12.5 Hz, blue; beta: 14–30.5 Hz, yellow; gamma:
5 33.5–100 Hz, magenta). Furthermore, we computed the relevance of each cluster per brain re-
6 gion by analyzing the amount of single subject trials during which a cluster was present. Group
7 clusters (Fig. 1, step 3) were traced back to single subject clusters and the amount of trials that
8 contributed to a single subject cluster (Fig. 1, step 2) was calculated and expressed as percent-
9 age. Percentages were averaged across subjects.

10 **Automatic Within-Group Classification**

11 A classifier was employed to test the specificity of region-specific spectral fingerprints. After
12 splitting each group into half (training and test group), group-level clusters were calculated for
13 the training group for all anatomical regions using k-means and GMM clustering. For each brain
14 region and participant of the test group, the similarity of spectral profiles was assessed as com-
15 pared to all brain regions of the 2nd-level group clusters of the training group by computing the
16 negative log-likelihood for all pairs of regions. This procedure, i.e., group assignment and clas-
17 sification, was repeated 1000 times (note that for the S-EO one subject was left out in every
18 iteration to yield an even number of participants in training and test groups). On each iteration,
19 an additional loop ($N = 100$) controlled for interindividual noise within a group by randomly
20 drawing the adequate number of subjects (i.e., $N_{S-EO} = 11$, $N_{S-BF} = 12$, $N_{CB} = 13$) from the group
21 with replacement, allowing a subject to enter multiple times or not at all. Put differently, within
22 one iteration ($N = 1000$) each participant belonged to either the training or the test group. To
23 account for individual differences, the group clusters were calculated 100 times with a different
24 subset chosen from the respective group each time and finally averaged to obtain a robust group

1 estimate. Based on the mode of clusters identified per brain region in the 2nd-level cluster anal-
2 ysis, the optimal number of clusters for the classification analysis was $k = 2$. Likelihood values
3 were ranked and averaged across iterations (20% trimmed mean). For further comparisons, only
4 corresponding ROIs (e.g., how is the Heschl ROI in the test set ranked based on the training set
5 Heschl ROI) were considered.

6 In addition to the descriptive report of the classification performance, here we tested whether a
7 specific ROI (of the test set) was classified significantly better by the corresponding area of the
8 training set as compared to all other 115 ROIs. This allowed us to exclude the possibility that
9 classification performance was caused by unspecific effects—that is, generic fingerprints. To
10 this end, each region’s mean rank (averaged across iterations) was tested against a distribution
11 of classification ranks generated from all other ROIs (null distribution).

12 **Automatic Cross-Group Classification**

13 In order to identify differences in region-specific spectral properties between the CB and S-BF,
14 we performed a cross-group classification. We employed the same classification procedure as
15 before; however, the classifier was trained on one group (S-BF), while the other (CB) was uti-
16 lized as the test set. As before, the classification procedure was repeated 1000 times, drawing a
17 subset of $N = 12$ per group on every iteration. The randomization of subjects chosen on each
18 iteration was identical to the one used for the within-group classification in the S-BF (this is the
19 reason why $N = 12$, instead of using all subjects of both groups). Thus, differences in the clas-
20 sification, as reflected by the ranks, could not be caused by the training set per se. In order to
21 understand whether some brain areas in the CB were not classified well based on the S-BF
22 spectral profiles (i.e., whether the classification of brain regions was different in the cross-group
23 condition as compared to the within-S-BF classification), we tested the cross-group classifica-
24 tion mean ranks against the distribution of ranks from the same ROI from the S-BF group (here
25 the null distribution). The distributions were generated by taking the classification rank of a

1 corresponding area from training and test set (i.e., Calcarine) across all iterations (see Fig. S2
2 for the distributions of all brain areas). We calculated the 95th percentile of the distribution and
3 tested whether the cross-group mean rank of the current region fell above (significant) or below
4 (not significant) this threshold.

5 To further assess the spectral profiles of brain areas that were significantly different in the cross-
6 group classification, post-hoc permutation statistics were applied to the raw, normalized region-
7 specific spectra (i.e., Fourier spectra without clustering procedure). This additional analysis was
8 performed to evaluate the cross-group differences between the sighted and the congenitally
9 blind groups by using a simpler, more established approach. The analysis confirmed the find-
10 ings from the spectral clustering approach. The spectral analysis was calculated as in the main
11 analysis (see above). For all significant brain regions separately, power was averaged across
12 voxels and segments, resulting in a single power value per frequency and per subject. Based on
13 frequency by subject matrices for the CB and the S-BF, group differences in spectral power
14 were tested against a distribution where the group assignment (CB vs. S-BF) was randomly
15 permuted ($N = 1000$). To control for multiple comparisons, we used FDR ($Q = 0.05$).

16 **Microstructural White Matter Properties**

17 DW-MRI data were acquired together with T1-weighted structural scans described above. We
18 used an echo planar imaging (EPI) sequence optimized for DWI-MRI of white matter covering
19 the whole brain (64 axial slices; bandwidth = 1502 Hz/Px, 104×128 matrix, TR, 8,200 ms; TE,
20 93 ms; flip angle, 90° ; slice thickness, 2 mm; voxel size, $2 \times 2 \times 2$ mm³). The protocol com-
21 prised three acquisitions yielding a total acquisition time of 9 min 51 s. This resulted in a total
22 of 120 diffusion-weighted volumes with six interleaved non-diffusion-weighted volumes (b
23 values of 1,500 s/mm²). DWI-MRI scans were acquired from a subset of the original sample of
24 16 blind and 12 sighted participants.

1 **DTI-MRI preprocessing and analysis.** Diffusion data processing initially corrected for eddy
2 current distortions and head motion by using FMRIB's Diffusion Toolbox (FDT; FMRIB Soft-
3 ware Library; FSL 5.0.1; <http://www.fmrib.ox.ac.uk/fsl/>; Jenkinson et al. 2012). For a more
4 accurate estimate of diffusion tensor orientations, the gradient matrix was rotated to correct for
5 head movement, using the `fdt_rotate_bvecs` program in FSL. We then used the Brain Extraction
6 Tool (Smith 2002) for brain extraction, also part of the FSL distribution. Analysis continued
7 with the reconstruction of the diffusion tensors using FSL's DTIFIT program. FA and RD maps
8 for each participant were calculated using the eigenvalues extracted from the diffusion tensors.
9 Note that FA maps are required in the early stages of TBSS to compute the registrations to MNI
10 standard space and subsequently create the diffusion skeletons. However, we focused our anal-
11 ysis on RD, as this is a more specific measure of diffusivity in white matter than FA or mean
12 diffusivity. Indeed, although several factors can contribute to producing particular RD values,
13 including the number of axons and axon packing and diameter, RD has been most consistently
14 related to myelin content along axons, with increased RD values reflecting higher demye-
15 lination (Song et al. 2002; Song et al. 2005; Klawiter et al. 2011; Zatorre et al. 2012). In animal
16 studies, directional measures such as RD, unlike summary parameters such as mean diffusivity
17 or FA, provide better structural details of the state of the axons and myelin (Aung et al. 2013).
18 Voxel-based analyses of RD maps were performed with TBSS (Smith et al. 2006). Participants'
19 FA maps (necessary to calculate the registrations to MNI standard space and create the RD
20 skeletons) were registered to the FMRIB58_FA template (MNI152 space and $1 \times 1 \times 1$ mm³)
21 using the nonlinear registration tool (Andersson et al. 2007). These registered FA maps were
22 first averaged to create a mean FA volume. A mean FA skeleton was then produced, represent-
23 ing the centers of all white matter tracts common to all participants in the study. Each partici-
24 pant's aligned FA data were then projected onto this skeleton by searching for the highest FA
25 value within a search space perpendicular to each voxel of the mean skeleton. This process was

1 repeated for the RD maps by applying the transformations previously calculated with the FA
2 maps. This resulted in individual RD skeletons for each participant. Finally, to assess white
3 matter differences between CB and sighted participants, independent-samples *t*-tests were per-
4 formed on the RD skeleton. Significant results are reported at FWE-corrected $p < 0.05$ using
5 threshold-free cluster enhancement (Smith and Nichols 2009) and a nonparametric permutation
6 test with 5,000 permutations (Nichols and Holmes 2002). Significant cluster results were aver-
7 aged and a mean value per participant, reflecting individual microstructural differences, was
8 obtained.

9 For the mapping between RD values and the standard probabilistic atlases of white matter path-
10 ways, all voxels that differed significantly in RD values between the CB and sighted were in-
11 cluded. We report only tracts that showed an overlap with these voxels, with the tracts from the
12 probabilistic atlas thresholded at 0.95 probability.

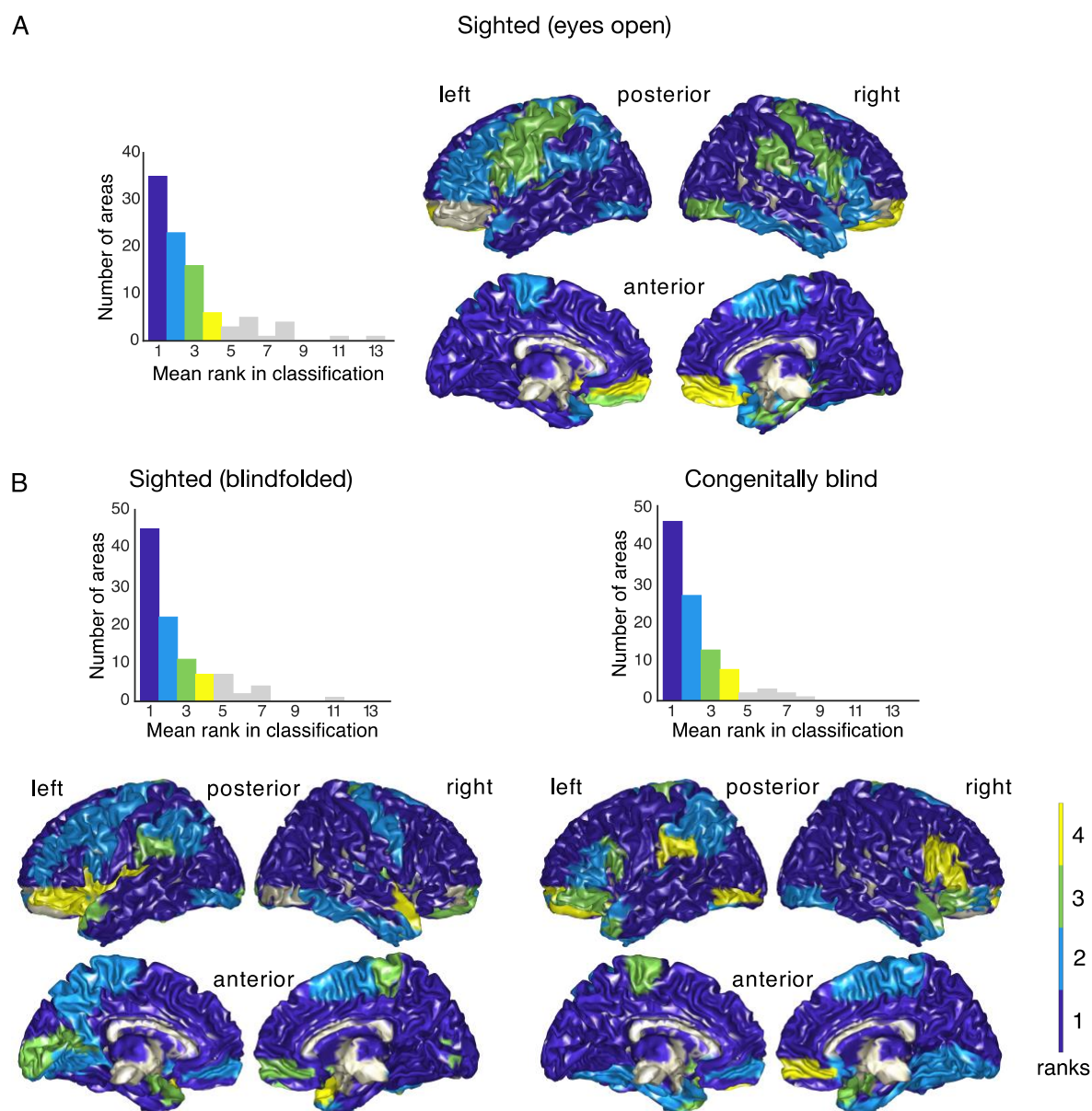
13 Spearman correlations were used to analyze the correlation between RD values (across groups)
14 and the spectral profiles of brain areas in the occipital cortex. For these areas, peak power values
15 were retrieved for all clusters (i.e., lines showing clear power peak) of the complex spectral
16 profile for each subject, and correlated with the RD values. Multiple comparisons were con-
17 trolled for by using FDR ($Q = 0.05$).

18 **Results**

19 **Spectral Fingerprints Replicate**

20 In our sample of sighted adults with open eyes we successfully replicated the classification of
21 individual brain regions by their spectral profiles as first reported by Keitel and Gross (2016).
22 Particularly, for all tested 115 ROIs the mean classification performance, indicated by the clas-
23 sification ranks, was high (Fig. 2A). Classification ranks refer to the probability of a region
24 being identified by the classifier: For example a mean rank of 1 indicates that a region was

1 correctly assigned (i.e., highest probability among all areas) on every iteration; a mean rank of
2 2 means that the assignment was correct in many but not all of the iterations (i.e., had the second
3 highest probability among all areas). Here, the mean rank (averaged across all iterations and
4 brain areas) obtained from the classifier analysis was 2.70 (range of ranks: 1–12.7, Keitel mean
5 rank = 1.8), or 2.32 when considering identification of the homologue (left/right hemisphere)
6 areas as a hit (Keitel homologue mean rank = 1.4). Mean ranks of all ROIs are depicted in the
7 histogram and surface plot in Fig. 2A. We statistically quantified the classification performance
8 using permutation tests. The mean classification rank of an area (e.g., right calcarine) was tested
9 against a distribution of classification ranks of all brain areas (except the current one, e.g., right
10 calcarine) accumulated across all iterations ($N = 1000$). For an area with a characteristic spectral
11 profile, classification between corresponding areas (e.g., right calcarine in training vs test set)
12 should be best and thus fall above the 95th percentile of the generated null-distribution. This
13 analysis revealed that for 97% of all areas classification was significantly better when identify-
14 ing themselves compared to all other regions.



1
 2 **Figure 2. Classification results for all experimental groups. (A) Sighted group with open eyes (replication**
 3 **sample).** *Left.* Histogram of mean ranks in classification across all 115 brain areas. 84.5% of the brain regions
 4 obtain a mean rank between 1 and 4, while 15.5% of regions were assigned ranks up to 13. (in Keitel and Gross
 5 (2016) all mean ranks were between 1 and 4). *Right.* Topography of mean ranks (colors match ranks from the
 6 histogram). **(B) Sighted group with eyes closed (left) and group of congenitally blind individuals (right).** For
 7 both groups, the histogram of mean ranks in classification across all 115 brain areas is displayed in the upper panel
 8 and the topography of mean ranks (colors match ranks from histogram) is displayed in the lower panel. Bin width
 9 is 1 for all subplots.
 10
 11 Furthermore—although the average optimal number of clusters per anatomical area was lower
 12 in our sample (3.4 ± 2.3 clusters per area (see Fig. 4 and Fig. S1, Supplementary Data) vs. 4.1
 13 ± 1.86 ($M + SD$) in Keitel and Gross (2016)—the clustering approach revealed comparable
 14 spectral fingerprints between the studies. Interestingly, for deeper subcortical brain structures

1 (e.g., thalamic and limbic areas), the clusters were less characteristic in the present data (i.e.,
2 only a few clusters per area with less specific shapes and high classification ranks; see Fig. S1),
3 possibly reflecting limitations of the signal-to-noise ratio of the used MEG system.

4 **Good Classification Within Sighted and Congenitally Blind Groups**

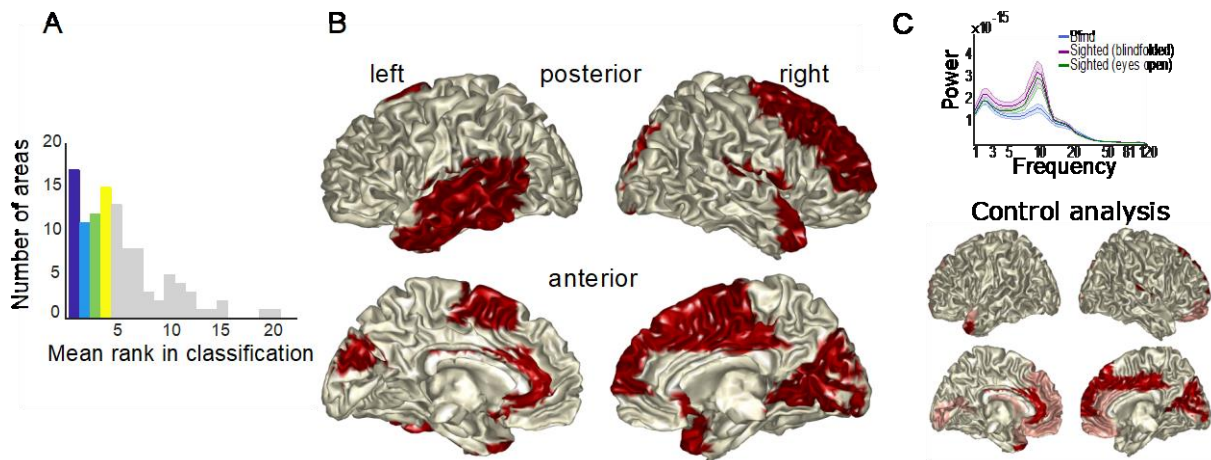
5 Our second hypothesis stated that, within a group of congenitally blind individuals, anatomical
6 areas are characterized by specific (although possibly altered as compared to the sighted) spec-
7 tral fingerprints. We performed the classification procedure for the CB and observed good clas-
8 sification ranks (similar to the ones observed for the S-EO; Fig. 2B; mean rank = 2.51, range =
9 1–10.3, homologue mean rank = 2.10, percent significant ROIs = 100%), indicating consistent
10 spectral clusters of brain areas in congenitally blind participants. The same procedure was per-
11 formed on the data of the S-BF and revealed similarly good classification ranks (S-BF: mean
12 rank = 2.64, range = 1–11.4, homologue mean rank = 2.17, percent significant ROIs = 98%) as
13 compared to the S-EO and CB.

14 Ensuring good within-group classification in the CB and the S-BF was an important prerequisite
15 for consecutive between-group analyses as it promised to prevent potential group differences
16 from arising from large within-group variance. Furthermore, the results showed a similar dis-
17 tribution of mean ranks across the cortical surface for both the CB and the S-BF group, such
18 that the majority of brain regions (~70 out of 115) was identified as best or second-best.

19 **Spectral Changes in Sensory and Right Frontal Regions in Congenitally Blind Individuals**

20 Based on the literature on intra- and cross-modal plasticity in the CB, we hypothesized that
21 spectral properties would differ between the congenitally blind and normally sighted individu-
22 als. To test if (and which) brain areas differed in their spectral properties between the two
23 groups, we implemented a cross-group classification drawing samples from the S-BF for the

1 training and samples from the CB for the test group. This analysis resulted in a mean rank of
2 5.3 (range = 1.09 - 27.17) (Fig. 3A).



3
4 **Figure 3. Cross-group comparison.** (A) Histogram of classification ranks. Bin width is 1. (B) The topographic
5 distribution of significantly different classification ranks in the cross-group classification is highlighted in red, as
6 tested by a permutation procedure. (C) The normalization spectrum used in the normalization procedure (i.e., the
7 average power spectrum across segments and voxels) is displayed, averaged across participants, separately for
8 each experimental group (upper panel). Shaded error bars reflect the standard error of the mean across participants.
9 Frequency is displayed at the x-axis, scaled logarithmically from 1 to 120 Hz; power on the y-axis. For visualization,
10 power was interpolated at 50 Hz to remove line noise. In order to test the impact of the normalization procedure
11 on our cross-group classification findings, a control analysis on the non-normalized data was performed
12 (lower panel). Confirming our findings, the control analysis revealed significant findings in various areas overlap-
13 ping with the areas observed in the main analysis (highlighted in dark red). However, we also observed several
14 additional areas with significant group differences (highlighted in light pink), as well as areas that, in contrast to
15 the main analysis, did not show significant group differences in the control analysis (see Table S2 for a full list).
16 Only findings at brain areas that showed group differences in both, the main cross-group and the control analyses
17 (i.e., with and without normalization procedure) will be considered for discussion.

18
19 In the cross-group analysis, 54.8% of 115 areas obtained classification ranks ranging from 1 to
20 4, while the automatic identification of the remaining regions was less precise (see Fig. 3A).
21 Beyond these descriptive procedures, we statistically compared classification results in the
22 cross-group condition as compared to the within S-BF group classification (see Fig. S2 for the
23 distributions of all brain areas). As seen in Fig. 3B, cross-group classification ranks were sig-
24 nificantly worse for several sensory as well as for right frontal areas, and an extended network
25 of left temporal brain regions. This suggests that spectral profiles in these brain regions were
26 different in the CB group as compared to the sighted (see Table 1 for all ROIs showing signif-

1 icantly worse classification compared to the null-distribution). A control analysis was per-
2 formed, to assure that group differences were not introduced by the normalization procedure
3 (i.e., reflecting differences in baseline power, Fig. 3C). While this analysis largely confirms our
4 findings, some of the brain areas did not show significant group differences in the control anal-
5 ysis and several additional areas with significantly worse classification ranks were found (see
6 Table S2 for a full list). Differences between the analyses are partly reflecting the advantages
7 of the normalization procedure, i.e., that it down-weights the neuronal activity (1/f power) that
8 is predominant across the brain and not specific to a given brain area (Fig. S5). In the following,
9 we focus on the interpretation of findings at ROIs that showed significant group differences in
10 both cross-group classification analyses (i.e., with and without normalization procedure). These
11 areas comprise: right Heschl’s gyrus, Calcarine gyrus, Cuneus, superior medial frontal gyrus,
12 middle cingulate cortex and left middle temporal pole and anterior cingulate cortex (highlighted
13 in Table 1).

14

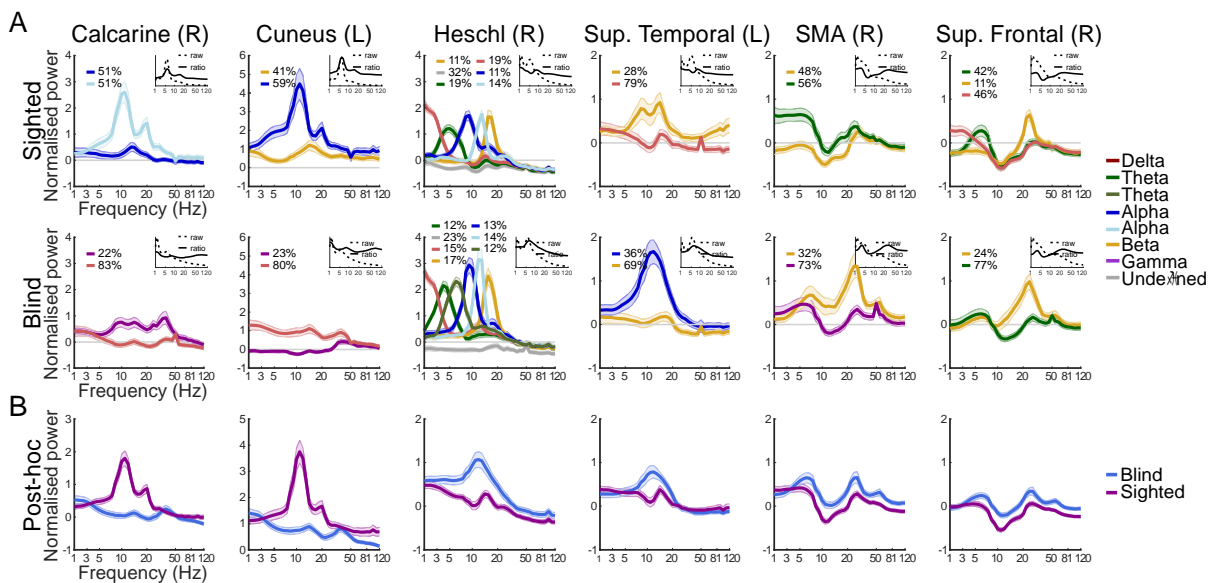
Coarse region	Hemisphere	
	Left	Right
Frontal		Superior frontal gyrus Middle frontal gyrus Superior medial frontal Rolandic Operculum
Visual	Cuneus	Calcarine gyrus Cuneus Superior occipital gyrus Lingual gyrus
Temporal	Superior temporal gyrus Middle temporal gyrus Middle temporal pole Inferior temporal gyrus	Heschl’s gyrus Middle temporal pole Superior temporal pole
Sensorimotor	Supplementary Motor Area	Supplementary Motor Area
Non-cortical	Olfactory Anterior cingulate cortex Cerebellum lobule VI Cerebellar vermis 3 Cerebellar vermis 9	Olfactory Middle cingulate cortex Cerebellum lobule X Cerebellar vermis 3 Cerebellar vermis 9

15 **Table 1. Table of all brain areas (out of 115) with significant classification differences.** Brain areas where
16 classification ranks were significantly different between the congenitally blind and the sighted (blindfolded) are
17 listed. The areas highlighted in bold showed significant group differences in the cross-group classification analyses

1 (i.e., with normalization procedure) and in the control analysis (i.e., without normalization procedure; see Fig. S5)
 2 and will be the focus of interpretation.

3

4 Interestingly, the brain areas identified as showing group differences in the spectral profiles in
 5 the cross-group classification were characterized by clusters comprising peaks with increased
 6 power at higher frequencies in the CB as compared to the S-BF participants (for an exemplary
 7 selection of brain areas with significant effects, see Fig. 4A; spectra of all brain areas are shown
 8 in Fig. S1). This pattern of results was observed for the auditory (with more power in the alpha
 9 and beta band in the CB as compared to the sighted participants) and the right frontal areas
 10 (more power in the theta band). In visual brain areas, power peaks were reduced in the alpha
 11 band for the CB as compared to the S-BF participants, in contrast, power was increased in the
 12 low-gamma band. Post-hoc permutation tests were performed to confirm these observations.
 13 To test differences in spectral signatures between the S-BF and CB, the raw Fourier spectra
 14 (i.e., without applying the spectral clustering) were extracted und subjected to permutation sta-
 15 tistics. Participants' group assignment (S-BF vs. CB) was permuted randomly (1000 permuta-
 16 tions) ($Q = 0.05$; false discovery rate [FDR] corrected p -value = .033; p -values < .033) (see Fig.
 17 4B and Fig. S3). Although the differences in low-gamma power in the calcarine between the
 18 CB and the S-BF were not significant in the post-hoc tests, see Fig. S6.



19

1 **Figure 4. Comparison of spectral profiles of sighted and blind participants.** (A) A selection of the spectral
2 profiles of the brain areas (columns) that showed significantly worse cross-group classification is displayed, sep-
3 arately for the S-BF and the CB (rows). Clusters are color-coded according to the frequency of the cluster ampli-
4 tude peak (legend on the right). The insets display the normalized mean power spectra (solid lines) and the unnor-
5 malized mean power spectra (dashed lines) (i.e., both without applying the spectral clustering). (B) Post-hoc anal-
6 yses of the spectral differences are displayed for the selection of brain regions (columns). Spectra represent nor-
7 malized mean power spectra (i.e., without applying the spectral clustering) at each ROI. Frequencies where power
8 differences were obtained between the groups are indicated by grey boxes (permutation test: $Q = 0.05$; FDR cor-
9 rected p -value = .033; p -values < .033). The groups are color-coded (legend on the right). In all panels: Shaded
10 lines indicate standard error of the mean.

11

12 **Spectral Changes Correlate with Structural Group Differences**

13 In order to test the relation of occipital spectral differences between the groups observed in the
14 classifier analyses and their relation to brain structure, we performed a diffusion-tensor imaging
15 (DTI) data analysis for a subsample of participants. The TBSS analysis revealed significantly
16 higher Radial Diffusivity (RD) values in a bi-hemispheric spatial cluster in posterior parts of
17 the brain (i.e., a cluster of voxels; displayed in Fig. S4B) for the CB as compared to the sighted
18 participants ($N_{sighted} = 12$, $N_{CB} = 16$; family-wise error (FWE)-corrected at the peak voxel,
19 two-sided $p = 0.05$; see Fig. S4B), indicating reduced white matter structural integrity in the
20 CB group. We used a probabilistic atlas of white matter pathways in MNI space (Thiebaut de
21 Schotten et al. 2011) to evaluate the overlap of the spatial cluster with known white matter
22 tracts. With the probabilistic atlas thresholded at 0.95, the TBSS cluster presents a significant
23 overlap with the posterior corpus callosum, the posterior inferior longitudinal fasciculus (bilat-
24 erally), the posterior inferior fronto-occipital fasciculus (bilaterally), and the optic radiations
25 (also bilaterally). This means that there is a 95% chance that the reduced white matter properties
26 identified in the CB group by the TBSS analysis primarily affect these white matter tracts.

27 Finally, RD values belonging to this spatial cluster were extracted and correlated with the power
28 of individual cluster peaks from the spectral profile of occipital brain areas (Spearman correla-
29 tions, FDR correction, $Q = 0.05$, corrected p -value = .0065, Fig. S4C). RD correlated negatively
30 with the alpha power in right ($\rho = -0.55$, $p = .005$) and left ($\rho = -0.81$, $p > 0.001$) calcarine,

1 as well as left cuneus ($\rho = -0.68, p > .001$), indicating that reduced white matter properties in
2 occipital areas were accompanied by reduced neuronal alpha power. Alpha and gamma power
3 of the remaining occipital regions did not correlate with the RD values (all p s $> .0065$; cf. Table
4 S1).

5 **Discussion**

6 Intrinsic brain rhythms likely are involved in various sensory, motor and higher-cognitive pro-
7 cesses. In congenitally blind individuals, behavioral performance in tasks involving the intact
8 senses has often been found to be enhanced (Lessard et al. 1998; Roeder et al. 2000; Roeder et
9 al. 2003; Gougoux et al. 2004; Hoetting et al. 2004; Roeder et al. 2004; Gougoux et al. 2005;
10 Hertrich et al. 2009; Dietrich et al. 2013), and structural and functional reorganization has been
11 observed in brain areas associated with the intact sensory systems (Pascual-Leone and Torres
12 1993; Sterr et al. 1998a; Sterr et al. 1998b; Roeder et al. 1999; Elbert et al. 2002; Stevens and
13 Weaver 2005; Gougoux et al. 2009; Schepers et al. 2012; Hertrich et al. 2013; Watkins et al.
14 2013), as well as in brain regions predominantly associated with the visual system (Elbert et al.
15 2002; Burton 2003; Roeder and Neville 2003; Noppeney et al. 2005; Pascual-Leone et al. 2005;
16 Noppeney 2007; Bedny et al. 2011; Voss and Zatorre 2012). Whether and how these behavioral
17 and neuronal adaptations affect brain rhythms outside of occipital brain regions had remained
18 unclear to date. Here, we identified spectral alterations in congenitally blind individuals aiming
19 at advancing a mechanistic understanding of adaptations to blindness. Furthermore, our find-
20 ings add to the ongoing debate regarding the functional relevance of brain rhythms. Confirming
21 our first two hypotheses, spectral clustering and classification procedures performed exception-
22 ally well for all groups (97–100% of areas were classified correctly in each group). This high-
23 lights consistent brain area-specific spectral properties across individuals within the sighted
24 group and, as demonstrated for the first time, within the congenitally blind group. Crucially,
25 cross-classification showed that visual deprivation co-occurred with changes in the spectral

1 profiles especially of sensory (auditory and visual), right-frontal, left middle temporal pole and
2 bilateral parts of the cingulate cortex. This was reflected in increased power in the theta-to-beta
3 frequency bands in the right primary auditory cortex and right-frontal brain regions. In visual
4 cortex, the sighted showed the alpha rhythm typically observed during rest, whereas in the con-
5 genitally blind individuals, clusters with expected decreased alpha power were accompanied by
6 a gamma (~40 Hz) peak, suggesting an altered inhibitory-excitatory balance in the blind. Addi-
7 tionally, a correlation analysis of individual spectral clusters in occipital cortex and microstruc-
8 tural white matter properties revealed a positive correlation between alpha (but not gamma)
9 power and white matter integrity in an extended bi-hemispheric spatial cluster in posterior parts
10 of the brain.

11 **Robust Classification of Brain Areas Based on Spectral Profiles**

12 Spectral profiles measured during resting state have revealed intrinsic brain rhythms that are
13 thought to be crucially involved in task-related processing (Smith et al. 2009; Hipp et al. 2012;
14 Engel et al. 2013; Raichle 2015; Keitel and Gross 2016; Sormaz et al. 2018). In the present
15 study, we replicated the observation that brain areas expressed characteristic spectral profiles
16 (Keitel and Gross 2016). Spectral clustering and automatic classification revealed spectral pro-
17 files, classification ranks, and distributions of classification ranks across the cortex in the
18 sighted (S-EO) group similar to the ones first reported by Keitel and Gross (2016). Spectral
19 profiles, for example of occipital regions, showed the typically observed alpha rhythm peaking
20 at ~10 Hz, which has repeatedly been found to be involved in attentional inhibition (Klimesch
21 et al. 2007; Buffalo et al. 2011; Jensen et al. 2012). Spectral peaks in the beta band (~20 Hz)
22 were prominent across frontal and central brain areas, resembling previously reported natural
23 frequencies of these brain areas (Rosanova et al. 2009; Ferrarelli et al. 2012; Keitel and Gross
24 2016), which may play a role in temporal motor processing (Fujioka et al. 2012; Arnal et al.
25 2015; Morillon and Baillet 2017).

1 While the spectral profiles of most brain areas resembled those reported by Keitel and Gross
2 (2016), for some brain areas the spectral profiles differed (see Fig. S1). In our study, deep sub-
3 cortical brain areas (in contrast to what has been reported by Keitel and Gross 2016) were not
4 classified well (Fig. S1). A possible reason for this is that the slightly shorter segment duration
5 used in the present study, which reduced the frequency resolution at lower frequencies, affected
6 the detection of low-frequency clusters. Alternately, a lower signal-to-noise ratio in deeper
7 brain areas in our data as compared to Keitel and Gross (2016) might be related to the employ-
8 ment of different MEG systems. Thus, possibly the recording system used and/or the sample of
9 participants tested potentially influenced the specific profiles of some brain areas more than
10 others. A test on a large dataset across different recording sites (i.e., several hundreds of record-
11 ings) will be necessary to clarify which spectral modes generalize across individuals of a larger
12 population. Importantly, within our sample, the spectral profiles were consistent across individ-
13 uals (i.e., only group clusters were reported in which at least ~70% of participants, and on av-
14 erage ~97% for the S-EO and ~94% for the S-BF group, contributed to each of the group-level
15 spectral clusters). In summary, our results highlight the robustness of brain area-specific spec-
16 tral profiles, suggesting that spectral profiles are characteristic properties reflecting the intrinsic
17 brain rhythms of cortical regions.

18 On the premise that changes in spectral profiles are functionally related to adaptive plasticity in
19 congenitally blind individuals, they are expected to be consistent (and not random) within a
20 group of congenitally blind individuals. A novel finding of our study which is in accord with
21 this hypothesis is that in the congenitally blind group, as with the sighted group, brain regions
22 were identified reliably based on their spectral clusters, suggesting spectral consistencies across
23 individuals (Fig. 2B, right; i.e., only group clusters were reported in which at least ~69% of
24 participants and on average ~95 % contributed to each spectral cluster). This result suggests

1 that adaptation of the cortex to visual deprivation leads to specifically and homogeneously al-
2 tered spectral fingerprints in the congenitally blind individuals.

3 **Selective Spectral Plasticity Across the Brain**

4 The main goal of our study was a comprehensive (in space and frequency), data-driven model
5 of spectral fingerprints in the congenitally blind as compared to sighted individuals in order to
6 provide insights into the neuronal mechanisms underlying plasticity in congenital blindness. In
7 the cross-group classification, brain areas of individual congenitally blind participants were
8 classified based on the group-level spectral clusters of the sighted (S-BF). In order to isolate
9 visual deprivation-related effects, the participant groups were well matched in our study (cf.
10 methods section). While in the cross-group classification the classification for the majority of
11 the brain areas was relatively good (i.e., low ranks; Fig. 3A), spectra related to sensory (auditory
12 and visual), right-frontal, left middle temporal pole and bilateral parts of the cingulate cortex
13 were classified significantly worse as compared to the within-sighted classification (Table 1;
14 Fig. 3B). These findings suggest that spectral properties may have been systematically altered
15 by visual deprivation-related plasticity for certain brain areas. Previously, a non-monotonic re-
16 lationship between plasticity and stability across the cortex has been reported using fMRI (Haak
17 and Beckmann 2019), whereas the plasticity observed in neuronal activity has been related to
18 gene expressions ([Ortiz-Terán et al. 2017](#)). Specifically, plasticity likely decreases from early
19 visual to mid-level cortex, but increases again further in the visual cortical hierarchy (Haak and
20 Beckmann 2019).

21 **Spectral Plasticity in Sensory Areas**

22 Altered spectral profiles in the congenitally blind individuals reflect changes in intrinsic brain
23 rhythms and might be linked to cognitive processes. Our findings highlight changes in spectral
24 properties of the auditory and visual cortices due to visual deprivation-related neuroplasticity.

1 The findings confirm previous reports demonstrating both cross-modal reorganization in visual
2 cortex (Burton 2003; Pascual-Leone et al. 2005; Bedny et al. 2011; Voss and Zatorre 2012;
3 Gudi-Mindermann et al. 2018; Rimmele et al. 2019) and intra-modal reorganization in auditory
4 cortex (Roeder and Neville 2003) in blind humans. Importantly, our findings extend these re-
5 sults by providing evidence for genuine changes in the processing mode of these regions, as
6 indicated by changes in the spectral characteristics, which contribute to a more mechanistic
7 understanding.

8 Visual brain areas classified as spectrally different between the sighted and the congenitally
9 blind individuals (in both the normalized and non-normalized control analysis) comprised right
10 primary visual cortex (calcarine sulcus) and cuneus (Table 1). In these areas, we observed in
11 the sighted two alpha-band clusters indicating distinct alpha brain rhythms. The posterior alpha
12 rhythm, which is typically observed in visual areas during resting state, likely reflects a mech-
13 anism of rhythmic inhibition of task-irrelevant neuronal circuits (Haegens et al. 2011). In our
14 data, we observed one cluster with a clear visual alpha peak at ~10 Hz and a smaller peak in the
15 beta-band (~20 Hz). A second alpha cluster was characterized by a smaller amplitude (note that
16 the two clusters are displayed by two separate lines in Fig. 4A), corroborating previous work
17 suggesting distinct visual alpha rhythms (Barzegaran et al. 2017). In contrast, in the congeni-
18 tally blind individuals these typical visual areas were characterized by a first cluster with a peak
19 at higher frequencies (low gamma, ~40 Hz) and a strongly reduced alpha power peak, as well
20 as a second cluster with low power in the alpha band (Fig. 4A, 4B). This observation is in line
21 with previous findings reporting a reduced or entirely absent alpha rhythm in the visual system
22 in blind individuals (Adrian and Matthews 1934; Noebels et al. 1978; Kriegseis et al. 2006;
23 Hawellek et al. 2013). The reduced alpha brain rhythm is thought to reflect atrophy or reorgan-
24 ization of the cortico-thalamic and cortical pathways, where posterior alpha generators have
25 been located (Lopes da Silva 1991; Lőrincz et al. 2009). It has been proposed that synchronized

1 posterior gamma activity is controlled by alpha (de-)synchronization and that it indicates in-
2 creased neuronal gain allowing feedforward processing of sensory information (van Kerkoerle
3 et al. 2014; Michalareas et al. 2016). The gamma rhythm in sighted individuals is typically
4 increased in the presence of a visual stimulus, but decreased or absent during rest. The gamma
5 rhythm may be preserved in blind individuals, as it evidently emerges locally in the occipital
6 cortex (Bastos et al. 2014; Marshall et al. 2018) being less impacted by atrophy of the thalamo-
7 cortical pathway. The presence of gamma activity in blind individuals during rest possibly re-
8 flects the alteration of the alpha-gamma excitation-inhibition balance, i.e., the reduced alpha
9 rhythm may result in a disinhibition of visual cortex increasing gamma band activity (Roeder
10 et al.). Functionally, while the reduced alpha rhythm in the congenitally blind may reflect cor-
11 tical and thalamo-cortical atrophy, the gamma-band rhythm might be related to the compensa-
12 tory reorganization of visual cortex, i.e., the recruitment of visual cortex during the processing
13 of non-visual tasks (Voss and Zatorre 2012; Striem-Amit et al. 2015; Bedny 2017; Voss 2019).
14 This conjecture is in line with our correlation analysis of the spectral profiles with microstruc-
15 tural white matter properties (Fig. S4, Table S1), which showed that only the alpha cluster cor-
16 related with the deteriorated white matter properties.

17 Additionally, we found altered spectral profiles in right auditory cortex (Heschl's gyrus; Table
18 1) with increases in power in specific frequency bands. In this area, we observed increased
19 power in the theta-to-beta frequencies in blind as compared to sighted participants (Fig. 4A,
20 4B). Importantly, in a comparison of auditory cortex spectral profiles during rest and during
21 speech comprehension, Keitel and Gross (2016) provided evidence for the functional relevance
22 of intrinsic delta, theta and beta brain rhythms, as observed during rest, for speech processing.
23 Speech tracking in the theta band (i.e., the phase-locking of auditory cortex oscillatory activity
24 to speech acoustics) is tightly linked to the temporal segmentation of syllables, as well as to the
25 prediction of upcoming stimuli (Giraud and Poeppel 2012; Gross et al. 2013; Haegens and Zion

1 Golumbic 2018; Rimmele, Gross, et al. 2018; Rimmele, Morillon, et al. 2018). Moreover, these
2 auditory cortex brain rhythms seem crucial for the segmentation of sound and music (Doelling
3 and Poeppel 2015; Morillon and Baillet 2017). Interestingly, previous research has shown that
4 congenitally blind individuals are able to comprehend ultra-fast speech at rates where speech
5 comprehension fails in sighted individuals (i.e., speech comprehension in sighted individuals
6 has been shown to fail at syllable rates faster than the theta range; Brungart et al. 2007; Ghitza
7 and Greenberg 2009). Furthermore, this ability to process ultra-fast speech in the congenitally
8 blind individuals has been linked to accelerated theta brain rhythms in right auditory cortex
9 (Hertrich et al. 2013) (see also: Trouvain 2007; Hertrich et al. 2009; Dietrich et al. 2013; Van
10 Ackeren et al. 2018). A possibility is that the altered spectral profiles in congenitally blind in-
11 dividuals reflect intrinsic brain rhythms that aid auditory temporal segmentation skills. Future
12 research, however, is required to confirm this speculation by relating the changes in brain
13 rhythms to behavioral performance.

14 **Spectral Plasticity Beyond Sensory Cortices**

15 Beyond spectral reorganization in sensory cortices, our data suggest that also right-hemispheric
16 frontal brain regions, bilateral parts of the cingulate cortex (left anterior, and right middle) and
17 the left middle temporal pole undergo adaptation. Spectral clusters of right medial superior
18 frontal gyrus and cingulate cortices were classified as different between the blind and the
19 sighted groups, with increased power in the theta-band in the blind group. Medial superior
20 frontal gyrus has been linked to various cognitive processes (Cohen et al. 2009; Cavanagh and
21 Frank 2014; Chander et al. 2016; Töllner et al. 2017), as well as theta-band brain rhythms (Ku-
22 bota et al. 2001; Onton et al. 2005; Cohen et al. 2008; Cohen 2011; Itthipuripat et al. 2013;
23 Hsieh and Ranganath 2014; Töllner et al. 2017; Eschmann et al. 2018; Başar-Eroglu et al.). For
24 instance, medial superior frontal gyrus has been implicated in cognitive control (Cavanagh and
25 Frank 2014), cognitive control of timing (Lewis and Miall 2003), decision making (Schmidt et

1 al. 2018), working memory (Itthipuripat et al. 2013; Hsieh and Ranganath 2014), temporal order
2 processing (Chander et al. 2016), or language processing (Binder et al. 2009). Some of these
3 processes are presumably altered in congenital blind individuals as compared to sighted indi-
4 viduals, such as language processing in a frontotemporal network (Roeder et al. 2002; Lane et
5 al. 2017), and temporal order processing (Roeder et al. 2004; Stevens and Weaver 2005).

6 The cingulate cortex (among other processes) has been associated with emotion, cognitive con-
7 trol and pain processing (Shackman et al. 2011). The left temporal pole, part of the anterior
8 temporal lobe, is thought to be involved in various processes including semantic (Binder et al.
9 1999; Visser and Lambon Ralph 2011) and conceptual (Baron and Osherson 2011) processing.
10 For both the cingular (Ortiz-Terán et al. 2016) and anterior temporal (Striem-Amit et al. 2018)
11 structures altered brain responses have been observed and proposed to be related to adaptive
12 behavior in congenitally blind as compared to sighted individuals. However, as the functionality
13 of these structures in congenital blindness, and particularly the role of endogenous brain
14 rhythms, remain little understood, we refrain from further interpreting these findings.

15 Note, that for some brain areas we found differences in spectral profiles in the cross-group
16 classification analysis (normalized data), however, not in the control analysis (on the non-nor-
17 malized data; Table 1; Fig. 3C, Table S2). We applied a previously established (Keitel & Gross,
18 2016) procedure. A disadvantage of the procedure is that it might introduce spurious activity
19 due to group differences in the normalization spectrum, as the procedure normalizes the spectra
20 across the brain. An advantage of the procedure is that it down-weights the neuronal activity
21 ($1/f$ power) that is predominant across the brain and not specific to a given brain area, which
22 otherwise complicates the spectral analysis. Thus, without the normalization the noise-level of
23 the data is increased due to individual differences in overall power across the spectrum and
24 across the brain. Although these arguments justify our choice, as precautionary measure we
25 only interpreted effects found in both analyses.

1 **Spectral profiles related to microstructural white matter properties**

2 A relevant question is to what extent our findings of altered spectral profiles of occipital cortex
3 in the congenitally blind are linked to structural differences. Our DTI analysis revealed com-
4 promised white matter integrity in congenitally blind individuals in visual association tracts
5 comprising the ventral visual stream. These tracts included the bilateral inferior fronto-occipital
6 fasciculus, connecting occipital and frontal brain areas and the bilateral inferior longitudinal
7 fasciculus, connecting occipital and anterior temporal cortices, which have been conjectured to
8 play a role in reading, writing and language semantics (Catani and Mesulam 2008); and the
9 bilateral optic radiations, linking the visual thalamus to the primary visual cortex. In addition,
10 white matter integrity was also compromised in the posterior corpus callosum, by which ho-
11 mologous visual cortices are interconnected (Restani and Caleo 2016). Reduced white matter
12 integrity in early (Shimony et al. 2006; Park et al. 2007), late (Wang et al. 2013; Hofstetter et
13 al. 2019), and congenitally blind individuals (Ptito et al. 2008; Wang et al. 2013; Aguirre et al.
14 2016; Reisleve et al. 2016), particularly in visual pathways (e.g., optic radiations and the corpus
15 callosum), is a well-documented finding. As myelination is proportional to the degree of neu-
16 ron activation during brain development (Demerens et al. 1996; Stevens et al. 2002; Fields
17 2004; Ishibashi et al. 2006; Gautier et al. 2015; Wake et al. 2015), the lack of visual processing
18 during development in congenital blindness likely causes the reduced white matter density in
19 visual pathways (Hill et al. 2014; Dietz et al. 2016; Restani and Caleo 2016; Anurova et al.
20 2019). Our findings show that reduced spectral power in the visual alpha rhythm co-occurred
21 with reduced white matter integrity in pathways that overlap with visual cortical pathways and
22 thalamo-cortical projections (i.e., negative correlation between alpha power and RD; Fig. S4B),
23 confirming previous accounts suggesting that reduced visual alpha power originates in struc-
24 tural impairment (Adrian and Matthews 1934; Noebels et al. 1978; Kriegseis et al. 2006;
25 Hawellek et al. 2013).

1 **Concluding Remarks**

2 The present study supports the findings of robust brain area-specific spectral profiles in the
3 human brain. We provide, for the first time, a whole-brain model of spectral fingerprints in
4 congenitally blind adults, and crucially, we were able to map spectral changes due to visual
5 deprivation to specific—namely right visual, auditory, frontal, left anterior temporal, and bilat-
6 eral cingulate—brain areas. We suggest that the power increases in the theta-to-beta frequency
7 bands in auditory and frontal brain regions may reflect adaptive sensory or higher cognitive
8 processing in blind individuals, while altered spectral profiles in visual brain regions (lower
9 alpha and a gamma peak) may indicate a change in the excitation-inhibition balance. While
10 these interpretations are tentative, the results pave the way for more targeted, task-based studies
11 that aim to identify the link between altered spectral profiles and adaptive behavior.

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18 **Competing Interests**

19 The authors declare that no competing interests exist.

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