Neuronal networks in children with continuous spikes and waves during slow sleep

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Epileptic encephalopathy with continuous spikes and waves during slow sleep is an age-related disorder characterized by the presence of interictal epileptiform discharges during at least >85% of sleep and cognitive deficits associated with this electroencephalography pattern. The pathophysiological mechanisms of continuous spikes and waves during slow sleep and neuropsychological deficits associated with this condition are still poorly understood. Here, we investigated the haemodynamic changes associated with epileptic activity using simultaneous acquisitions of electroencephalography and functional magnetic resonance imaging in 12 children with symptomatic and cryptogenic continuous spikes and waves during slow sleep. We compared the results of magnetic resonance to electric source analysis carried out using a distributed linear inverse solution at two time points of the averaged epileptic spike. All patients demonstrated highly significant spike-related positive (activations) and negative (deactivations) blood oxygenation-level-dependent changes (P < 0.05, family-wise error corrected). The activations involved bilateral perisylvian region and cingulate gyrus in all cases, bilateral frontal cortex in five, bilateral parietal cortex in one and thalamus in five cases. Electrical source analysis demonstrated a similar involvement of the perisylvian brain regions in all patients, independent of the area of spike generation. The spike-related deactivations were found in structures of the default mode network (precuneus, parietal cortex and medial frontal cortex) in all patients and in caudate nucleus in four. Group analyses emphasized the described individual differences. Despite aetiological heterogeneity, patients with continuous spikes and waves during slow sleep were characterized by activation of the similar neuronal network: perisylvian region, insula and cingulate gyrus. Comparison with the electrical source analysis results suggests that the activations correspond to both initiation and propagation pathways. The deactivations in structures of the default mode network are consistent with the concept of epileptiform activity impacting on normal brain function by inducing repetitive interruptions of neurophysiological function.

Keywords: continuous spikes and waves during slow sleep (CSWS); EEG-fMRI; EEG source analysis; perisylvian region; remote inhibition; children
Introduction

Epileptic encephalopathy with continuous spikes and waves during slow sleep (CSWS) is an age-related reversible disorder characterized by acquired variable neuropsychological impairment, epilepsy with heterogeneous seizure types, and the presence of the interictal electroencephalographic (EEG) findings of intense sub-continuous paroxysmal activity consisting of spike-wave complexes that usually occupy more than 85% of non-rapid eye movement (REM) sleep (Tassinari et al., 2005). This disorder may be attributed to different aetiologies (symptomatic cases with various lesions, cryptogenic epilepsies and even idiopathic cases, for example, in the form of Landau–Kleffner syndrome or atypical benign partial epilepsy of childhood). CSWS is usually associated with manifold psychomotor and cognitive deficits (auditory agnosia, acquired aphasia, dysfunctions of the frontal lobe and short-term memory deficits, pseudo-bulbar palsy, global mental deterioration, impaired spatial orientation, apraxia and hemineglect, psychotic states and autistic features, attention deficit and hyperactivity and aggressiveness) (Roulet Perez et al., 1993; Maquet et al., 1995; Shafir and Prensly, 1995; Galanopoulou et al., 2000; Eriksson et al., 2003; Scholtes et al., 2005; Deona and Roulet, 2006). Despite this heterogeneity, however, most patients with CSWS demonstrate common features: substantial increases in paroxysmal activity during sleep and a complex morphology of interictal epileptiform discharges with bilateral and diffuse spikes and waves, which tend to generalize (Tassinari et al., 2005; Nickels and Willems, 2008). Although in the first description of the epileptic encephalopathy with CSWS by Patry et al. (1971), the authors suggested a ‘particularly active synchronizing system’ that ‘could account for extreme activation of the spike and wave spikes’ during sleep, and different studies have provided arguments in favour of secondary bilateral synchrony (Gaggero et al., 1995; Maquet et al., 1995; Morrell et al., 1995; De Tiege et al., 2004; Luat et al., 2005), understanding of the underlying mechanisms is very limited.

Simultaneous recording of EEG and blood-oxygenation level-dependent (BOLD) functional magnetic resonance imaging (EEG-fMRI) is a non-invasive technique that allows evaluation of haemodynamic changes in the brain correlated with pathological activity on the scalp EEG acquired within the scanner (Hamandi et al., 2005; Gotman et al., 2006; Laufs and Duncan, 2007). Due to whole-brain coverage and relatively high spatial resolution, this technique is able to characterize neuronal networks associated with EEG patterns in a unique way (Vulliemoz et al., 2009). BOLD patterns reflecting networks have been demonstrated in relation to primary and secondary generalized epileptiform activity (Aghakhani et al., 2004; Hamandi et al., 2006; Moeller et al., 2008) and hypsarrhythmia (Siniatchkin et al., 2007b). In focal epilepsies, the EEG-fMRI has revealed not only neurometabolic changes in brain regions corresponding to the spike localization, but also in functionally significant brain areas remote of epileptic activity (Kobayashi et al., 2006; Jacobs et al., 2007, 2009; Laufs et al., 2007). The addition of electrical source imaging may allow the identification of areas of spike generation and propagation and therefore enhance the interpretation of fMRI results (Boor et al., 2007; Grova et al., 2008; Groening et al., 2009; Vulliemoz et al., 2009, 2010a).

Here, we characterized the neuronal networks associated with epileptic seizures using simultaneous EEG-fMRI recordings in children with epileptic encephalopathy and CSWS during drug-induced sleep. The study aims to distinguish brain areas which are common across patients and are involved in the pathological process underlying CSWS.

Materials and methods

Subjects

Between May 2005 and June 2009, patients with CSWS were recruited in the Northern German Epilepsy Centre, Raisdorf and in the University Hospital of Paediatric Neurology, Kiel, Germany. The inclusion criteria were: (i) the presence of epileptic activity during >85% of non-REM sleep; (ii) new diagnosis of epileptic encephalopathy with CSWS before specific treatment of this condition; and (iii) bilateral and diffuse EEG abnormalities (Fig. 1). Diagnosis was made according to the International League against Epilepsy 2001 classification scheme (Commission on Classification and Terminology of the International League against Epilepsy, 2001). Eight boys and four girls with CSWS, aged between 3 and 13 (mean 8.5 ± 2.8 years) at the time of recording, met the criteria. The clinical and demographic characteristics of the group are shown in Table 1. All patients were investigated in the neuroimaging laboratory of the University Hospital of Paediatric Neurology, Kiel.

The neurological examination and structural MRI (high-resolution T1-, T2-, fluid attenuated inversion recovery T2- and diffusion-weighted imaging) were performed before inclusion in the study. Routine sleep EEGs under sedation (32 electrodes according to the International 10–20 System, the same sedation as for EEG-fMRI, i.e. oral chloral hydrate 75 mg/kg) were recorded 1–2 days before the EEG-fMRI investigation and were evaluated by at least two specialists independently, who confirmed the presence of CSWS. All patients presented with neuropsychological abnormalities: global retardation, aphasia, attention-deficit hyperactivity disorder, conduct disorder and autism (Table 1). None of the children suffered from Landau–Kleffner syndrome. All children were sedated 30 min before MRI scanning and the EEG-fMRI recordings were performed when the children were asleep. Foam pads were used to help secure the EEG leads, minimize motion and improve patient comfort. A paediatrician was present throughout the examination. The study was performed according to

Abbreviations: BOLD = blood oxygenation dependent; CSWS = continuous spikes and waves during slow sleep; fMRI = functional magnetic resonance imaging; MNI = Montreal Neurological Institute; REM = rapid eye movement; SPM = statistical parametric mapping
Figure 1: Typical pattern of spike-and-wave paroxysms in a patient with CSWS. The example represents the EEG obtained inside the scanner (original montage).

Table 1: Clinical and demographic characteristics of patients with CSWS and observed pathological activity on EEG during 20 min fMRI acquisition

<table>
<thead>
<tr>
<th>Patient</th>
<th>Sex/age/age of epilepsy onset</th>
<th>Seizure type</th>
<th>MRI</th>
<th>Antiepileptic drugs</th>
<th>Neuropsychology</th>
<th>Number of spikes during fMRI</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>F/8/4</td>
<td>CPS, GTCS</td>
<td>–</td>
<td>VPA</td>
<td>Aphasia, ADHD</td>
<td>1375</td>
</tr>
<tr>
<td>2</td>
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<td>–</td>
<td>PVL</td>
<td>VPA</td>
<td>Global retardation</td>
<td>1656</td>
</tr>
<tr>
<td>3</td>
<td>M/9/3</td>
<td>SPS, GTCS</td>
<td>PVL</td>
<td>STM, VPA</td>
<td>Global retardation</td>
<td>1898</td>
</tr>
<tr>
<td>4</td>
<td>M/9/7</td>
<td>CPS, GTCS</td>
<td>Left HM</td>
<td>VPA, LTG</td>
<td>Aphasia, CD</td>
<td>1738</td>
</tr>
<tr>
<td>5</td>
<td>F/13/6</td>
<td>CPS, GTCS</td>
<td>–</td>
<td>VPA, LTG</td>
<td>Global retardation</td>
<td>1566</td>
</tr>
<tr>
<td>6</td>
<td>M/7/5</td>
<td>CPS</td>
<td>PVL</td>
<td>VPA, LEV</td>
<td>Global retardation</td>
<td>2308</td>
</tr>
<tr>
<td>7</td>
<td>F/10/4</td>
<td>CPS</td>
<td>–</td>
<td>VPA, TPM</td>
<td>Global retardation</td>
<td>2407</td>
</tr>
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<td>8</td>
<td>M/9/4</td>
<td>–</td>
<td>–</td>
<td>VPA</td>
<td>Aphasia, ADHD, CD</td>
<td>1398</td>
</tr>
<tr>
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<td>–</td>
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<td>Cortisone</td>
<td>Global retardation</td>
<td>1434</td>
</tr>
<tr>
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<td>–</td>
<td>VPA</td>
<td>Global retardation</td>
<td>1163</td>
</tr>
<tr>
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<td>–</td>
<td>–</td>
<td>VPA</td>
<td>Autism</td>
<td>1315</td>
</tr>
<tr>
<td>12</td>
<td>M/3/1</td>
<td>CPS</td>
<td>PVL</td>
<td>LEV/TPM</td>
<td>Global retardation</td>
<td>1272</td>
</tr>
</tbody>
</table>

ADHD = attention deficit hyperactivity disorder; CD = conduct disorder; CPS = complex partial seizures; GTCS = generalized tonic-clonic seizures; HM = hippocampal malrotation; LEV = levetiracetam; LTG = lamotrigine; PVL = periventricular leucomalacia; SFS = simple focal seizures; STM = sulthiame; TPM = topiramate; VPA = valproate.

Age is given in years. SPIKE represent frequency in 20 min recording. * - negative MRI.
the Declaration of Helsinki and was approved by the local ethics committee. The parents gave written informed consent.

Data acquisition

The EEG was continuously recorded inside the MRI scanner from 32 scalp sites (10–20 system plus FC1, FC2, CP1, CP2, FC5, FC6, CP5, CP6, TP9, TP10, FT9, FT10) with a reference located between Fz and Cz. Sintered Ag/AgCl ring electrodes were attached using the ‘BrainCap’ (Falk-Minow Services, Hersching-Breitbrunn, Germany), which is part of the magnetic resonance-compatible EEG recording system ‘BrainAmp-MR’ (Brainproducts Co., Munich, Germany). Electrode impedance was kept below 7 kΩ (after subtraction of the value of the safety resistors). Data were transmitted from the high-input impedance amplifier (250-Hz low-pass filter, 10-s time constant, 16-bit resolution, dynamic range 16–38 mV) which was placed directly behind the head coil inside the scanner room and connected to a computer located outside the scanner room via a fibre optic cable. The scanner (10-MHz sampling rate) was synchronized with the EEG amplifier (5-kHz sampling rate). Online correction of gradient artefacts based on the averaged artefact subtraction algorithm was performed using RecView software (Brainproducts Co., Munich, Germany) enabling visual inspection of spikes throughout the recording.

BOLD-sensitive MRI was performed with a 3 Tesla MRI scanner (Philips Achieva, Philips, Best, The Netherlands) and a standard, 8-channel sensitivity encoding head coil. A single-shot T$_2^*$-weighted gradient-echo planar imaging sequence (repetition time = 2250 ms, echo time = 45 ms, 30 slices, 64 × 64 matrix, slice thickness = 3.5 mm, field of view = 200 mm, flip angle = 90°). Data from 540 brain volumes were acquired during the 20-min fMRI session. The first five images were discarded to ensure steady-state longitudinal magnetization. An anatomical MRI for superimposition with functional images was acquired using a T$_1$-weighted three-dimensional (3D) multi planar reformating sequence (1 mm slice thickness, 208 × 208 matrix, 150 slices, field of view = 208 mm, echo time = 3.6 ms, repetition time = 7.8 ms, flip angle = 8°, number of signal averages = 2).

EEG processing

Gradient artefacts as a result of electromagnetic distortion of the EEG through static and dynamic magnetic field during magnetic resonance data acquisition were removed offline using the averaged artefact subtraction method as implemented in the BrainVision Analyser 1.05 software (Brainproducts Co., Munich, Germany) (Allen et al. 1998, 2000). After artefact correction, the data were down-sampled to 250 Hz and filtered at 0.03–75 Hz. Heartbeat-related artefacts were reduced by means of the multiple source correction algorithm as implemented in Brain Electrical Source Analysis software BESA 5.2 (MEGIS Software Co., Munich, Germany) and described in Sinitchkin et al. (2007a). Epileptiform activity was identified semi-automatically using BESA, following the method of Bast et al. (2004). Firstly, the EEG was inspected to select a typical spike, showing similar morphology and topography to those marked on the EEG acquired outside the scanner (routine EEGs from the Northern German Epilepsy Centre). A spatiotemporal pattern-matching algorithm combined with visual inspection was then used to identify similar spikes throughout the recording (Scherg et al., 1999, 2002). All spikes selected had similar shape and topography. The spike selection based on spike topography may enhance sensitivity of EEG-fMRI studies in epilepsy (Sinitchkin et al., 2007a; Vulliemoz et al., 2010). The quality of spike detection was confirmed by two experienced neurophysiologists independently, resulting in a consensus concerning the spike set to be used for fMRI analysis and EEG source localization.

EEG source analysis

Following application of averaged artefact subtraction and multiple source correction methods, data quality was sufficient to allow reliable spike identification and electrical source imaging (Sinitchkin et al., 2007a). Because spikes in patients with CSWS occurred in long sequences and bursts, one could suggest that spikes in the sequence may be related to propagated epileptic activity and the area of the initial activity (generator) of the spike within the sequence may be different than the area of initial activity of the first spike in the sequence (Ebersole et al., 2000). Indeed, a preliminary analysis of all averaged spikes revealed a poor correspondence between electrical source imaging and fMRI results, especially for the initial epileptic activity. Therefore, for the purpose of electrical source imaging, we selected the first spike in every run of spikes preceded by a sufficiently long period of normal background (at least 2 s). The selected spikes were averaged and aligned to the global field power peak and projected in the epoch of ±500 ms around this peak (Michel et al., 2004a). As suggested by Ebersole (2000), we checked whether different voltage topographies appear over the course of the spike, indicating propagation. Two time frames were chosen to characterize spike topography: (i) from the beginning of the spike to the time point corresponding to 50% of the rising phase as an epoch characterizing a source of the possible spike generator: t = t$_{spike\_onset}$ (see Scherg et al., 1999; Huppertz et al., 2001; Lantz et al., 2003a); and (ii) an epoch around the peak of the spike: t = t$_{spike\_propagation}$. The precise timings of the frames were local minima in the time course of the dissimilarity index, which is a measure inversely related to the spatial correlation between two scalp voltage map topographies: a minimum of dissimilarity therefore reflects a period of map stability (Lantz et al., 2003a; Michel et al., 2004b) (Figs 2, 3 and 4, parts C and D). Source analysis at t = t$_{spike\_onset}$ and t = t$_{spike\_propagation}$ was performed using a distributed linear inverse solution, based on a local autoregressive average model of the current density of the brain using the Cartool software (http://brainmapping.unige.ch/Cartool.htm), giving the two solutions: electrical source imaging-o and electrical source imaging-p, respectively (Grave de Peralta Menendez et al., 2001; Michel et al., 2004b). Validation and application of this localization method in epileptic EEG has been shown in previous studies (Brodbbeck et al., 2009, 2010; Groening et al., 2009; Vulliemoz et al., 2009, 2010b).

Local autoregressive average was calculated in a simplified realistic head model called SMAC (Spinelli et al., 2000); the brain surface is extracted from the individual MRI and the best fitting sphere is estimated. Then the MRI is warped according to the ratio of the sphere radius and the real surface radius. Depending on brain size, between 3500 and 4500 solution points were defined in regular distances within the grey matter. The forward problem is then solved with an analytical solution using a realistic head model. Additional details can be found in Groening et al. (2009).

To assess the presence of population-wide effects, the results of the single-subject source analyses were then taken to a second level group analysis (Friston et al., 1999). The results of the local autoregressive average source reconstruction were transformed to the standard Montreal Neurological Institute (MNI) space using Cartool and saved as a volume. The volumes were realigned, smoothed with the isotropic 9 mm Gaussian kernel and spatially normalized to the MNI template using SPM5 software (Statistical Parametric Mapping, Wellcome Department of Imaging Neurosciences, UCL, UK, http://www.fil.ion
Functional MRI processing

The fMRI data were analysed using SPM5. All volumes were realigned to the first volume and spatially normalized to the MNI template of the SPM software. Images were then smoothed using an isotropic Gaussian kernel of 6 mm and high-pass filtered at a cut-off of 128 s. The pre-processed fMRI time series were statistically analysed at an individual level using the General Linear Modelling approach, in which each spike was treated as a single event. Each event was represented as a stick function, convolved with a canonical haemodynamic response function (peak at 6 s relative to onset, delay of undershoot = 16 s, length of kernel = 32 s) as implemented in SPM5 (Friston et al., 1995). In each individual, one-tailed t-tests were applied to test for regional spike-related BOLD increases or decreases. At the voxel level, the significance level was set at $P < 0.05$ after correction for multiple comparisons across the whole brain using the family-wise error correction method as implemented in SPM5 (Friston et al., 1995). This corresponded to t-values above 4.7. An extent threshold of five contiguous voxels was also applied. Individual statistical parametric t-maps were colour-coded and superimposed onto the individual coregistered T1-weighted images. Using the parameter estimates obtained by single-subject analyses, we performed a second-level random effect group analysis (one sample t-test) to test for
typical BOLD signal changes at the population level. The threshold was set at $P < 0.05$ (false discovery rate corrected). The resulting statistical maps were displayed in MNI space.

**Results**

**EEG-functional MRI study**

All patients demonstrated highly significant positive and negative spike-related BOLD signal changes ($P < 0.05$ corrected; Figs 2–4 and Table 2). Despite high spike frequency, the time series of BOLD responses were characterized by sufficient variability and visual comparison of the BOLD time series in the regions of significantly correlated change and EEG showed a good correspondence with the epileptic spikes (Fig. 5). Maximal $t$-values varied between 5.9–19.7 for BOLD increases and between 4.7 and 25.4 for BOLD decreases. BOLD increases were observed bilaterally in the perisylvian region (insula, superior temporal gyrus, inferior frontal gyrus) and in the anterior cingulate gyrus/prefrontal cortex in all patients, the thalamus in five (42%), the temporal lobe in three (25%), the frontal cortex in five (42%) and in the parietal and occipital cortices in one (8%). BOLD decreases were revealed bilaterally in the parietal cortex in 11 patients (92%), in the precuneus in 10 (83%), in the medial prefrontal and frontal cortices in five (42%) and in the caudate nucleus in four (33%). The localization of maximal $t$-values for both positive and negative BOLD effects varied from patient to patient. No systematic difference in patterns was apparent between symptomatic and cryptogenic cases. The structural abnormalities found in six patients...
The group analysis revealed a significant bilateral positive BOLD effect in the perisylvian region, cingulate and prefrontal cortices and thalamus (Fig. 6 and Table 3), in a pattern resembling an arch. BOLD decreases were observed in the precuneus and bilaterally in the parietal cortex and in the caudate nucleus.

**EEG source analysis**

The results of the single-subject EEG source analysis were broadly similar across patients for both time points (Table 4). The initial epileptic activity was localized in the frontal cortex in three patients (25%), the perisylvian region/insula in eight (67%) and the parietal and occipital cortices in one (8%). A good correspondence between the initial epileptic activity as revealed by the EEG source analysis and brain areas with regions of BOLD increase was found in 11 patients (92%). The epileptic activity propagated bilaterally in the perisylvian region in all patients, the frontal cortex in nine (75%), the posterior brain areas in eight (67%) and the temporal lobe in seven (58%). The source analysis for the propagation time point revealed source activity in the perisylvian region with corresponding spike-related BOLD increases in all patients. In six patients (50%), there was good correspondence between propagated source activity in the frontal and temporal cortices and regions of BOLD increase. Propagated source activity in the parietal cortex was in agreement with BOLD decreases in three patients (25%).

The group analysis of EEG source activity did not reveal any constant source for the rising phase of the spike, possibly because of different individual localization of the initial epileptic activity (Fig. 7A and Table 4). However, the group analysis of the
Table 2  Results of EEG-fMRI analyses for positive and negative CSWS-related BOLD effects

<table>
<thead>
<tr>
<th>t-value</th>
<th>Persylvian insula</th>
<th>Cingulate prefrontal</th>
<th>Temporal cortex</th>
<th>Thalamus</th>
<th>Frontal parietal</th>
<th>t-value</th>
<th>Precuneus</th>
<th>Parietal cortex</th>
<th>Medial prefrontal</th>
<th>Caudate nuclei</th>
<th>Occipital</th>
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<td>1</td>
<td>19.69 (T)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>25.39 (CN)</td>
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<td>+</td>
<td>+</td>
<td>+</td>
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</tr>
<tr>
<td>2</td>
<td>7.74 (TC)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>13.63 (CN)</td>
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<td>+</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>6.14 (PC)</td>
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<td>+</td>
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<td>+</td>
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<td>+</td>
<td>9.74 (FC)</td>
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<td>+</td>
<td>+</td>
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<td>+</td>
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<td>+</td>
<td>10.05 (OC)</td>
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</table>

All t-values represent global maximum and are P < 0.05, corrected. In all patients, only bilateral changes were observed. In = insula; CG = cingulate gyrus; FC = frontal cortex; PC = parietal cortex; TC = temporal cortex; OC = occipital cortex; P = precuneus; T = thalamus; CN = caudate nucleus.

Figure 5  Positive BOLD response (A) that was obtained in Patient 1 in a voxel with the maximal t-value in the right insular region, which was chosen because of the correspondence between fMRI results and results of the electrical source imaging of the rising phase of the spike (beginning of epileptic activity). The first 70 s of recording are enlarged (B) to show the relation between variation of the fitted positive BOLD response and spike frequency. Note that frequent spikes were associated with an increase in BOLD response and gaps between spikes corresponded well with BOLD decreases. This good correlation between the occurrence of spikes and BOLD signal changes explains highly significant results of EEG-fMRI studies in patients with CSWS.
propagated epileptic activity (peak) was able to describe electrical sources in the left and right perisylvian region, left and right temporal cortices (mostly in the superior temporal gyrus), temporoparietal junction and the left prefrontal and parietal cortices.

The results of the source analysis on the group level show good correspondence with results of the random effect group analysis of fMRI data, especially with activation in the perisylvian region and deactivation in the parietal cortex and temporoparietal junction. This positive correspondence between fMRI results and EEG pattern supports the contention that fMRI is able to reveal a physiologically meaningful pathological network.

**Discussion**

This study revealed the following main findings: (i) independent of aetiology, a common pattern of CSWS-related BOLD increases bilaterally in the perisylvian region; (ii) this pattern was confirmed using EEG source reconstruction showing a common bilateral pattern of propagation in the perisylvian region associated with different individual localization of the initial epileptic activity; (iii) there is frequent involvement of the prefrontal cortex and cingulate gyrus in both EEG-fMRI and source analysis; (iv) common BOLD increases in the thalamus and BOLD decreases in the caudate nuclei are associated with CSWS, especially revealed by the group analysis; and (v) there is a common pattern of BOLD decreases in the default mode network.

Table 3  Results of the group analysis (maximum of activation in the cluster)

<table>
<thead>
<tr>
<th>Structures</th>
<th>t-value</th>
<th>Voxels</th>
<th>MNI coordinates</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>BOLD increases</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Right perisylvian region and insula</td>
<td>7.27</td>
<td>293</td>
<td>36/0/-9</td>
</tr>
<tr>
<td>Left perisylvian region and insula</td>
<td>6.69</td>
<td>143</td>
<td>-51/-6/15</td>
</tr>
<tr>
<td>Right prefrontal cortex and cingulate gyrus</td>
<td>7.80</td>
<td>252</td>
<td>10/-9/51</td>
</tr>
<tr>
<td>Left prefrontal cortex and cingulate gyrus</td>
<td>6.19</td>
<td>154</td>
<td>-39/-15/54</td>
</tr>
<tr>
<td>Thalamus</td>
<td>4.11</td>
<td>54</td>
<td>12/-12/6</td>
</tr>
<tr>
<td><strong>BOLD decreases</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Precuneus</td>
<td>4.40</td>
<td>271</td>
<td>5/-56/30</td>
</tr>
<tr>
<td>Right parietal cortex</td>
<td>4.40</td>
<td>364</td>
<td>51/-60/27</td>
</tr>
<tr>
<td>Left parietal cortex</td>
<td>4.54</td>
<td>117</td>
<td>-51/-69/30</td>
</tr>
<tr>
<td>Caudate nucleus</td>
<td>4.22</td>
<td>45</td>
<td>-8/15/5</td>
</tr>
</tbody>
</table>

Figure 6  Results of the random effect group analysis for both positive and negative BOLD signal changes (false discovery rate corrected, \( P < 0.05 \)).
Figure 7 Results of the source analysis are presented for the initial epileptic activity on the individual level (A) and for the propagated epileptic activity on the group level (B).

Table 4 Results of the EEG source analysis for the initial and propagated epileptic activity

<table>
<thead>
<tr>
<th>Patient number</th>
<th>Spike number</th>
<th>Initial activity</th>
<th>Propagated activity</th>
<th>Correspond</th>
<th>Initial/propagated</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Perisylvian insula</td>
<td>Frontal</td>
<td>Parietal occipital</td>
<td>Perisylvian insula</td>
</tr>
<tr>
<td>1</td>
<td>54</td>
<td>R</td>
<td>B</td>
<td>B</td>
<td>B</td>
</tr>
<tr>
<td>2</td>
<td>24</td>
<td>L</td>
<td>B</td>
<td>L</td>
<td>B</td>
</tr>
<tr>
<td>3</td>
<td>88</td>
<td>R</td>
<td>B</td>
<td>L</td>
<td>L</td>
</tr>
<tr>
<td>4</td>
<td>91</td>
<td>L</td>
<td>B</td>
<td>B</td>
<td>B</td>
</tr>
<tr>
<td>5</td>
<td>43</td>
<td>L</td>
<td>B</td>
<td>B</td>
<td>B</td>
</tr>
<tr>
<td>6</td>
<td>47</td>
<td>L</td>
<td>B</td>
<td>B</td>
<td>B</td>
</tr>
<tr>
<td>7</td>
<td>26</td>
<td>L</td>
<td>B</td>
<td>L</td>
<td>L</td>
</tr>
<tr>
<td>8</td>
<td>70</td>
<td>L</td>
<td>B</td>
<td>B</td>
<td>B</td>
</tr>
<tr>
<td>9</td>
<td>56</td>
<td>L</td>
<td>B</td>
<td>B</td>
<td>R</td>
</tr>
<tr>
<td>10</td>
<td>23</td>
<td>L</td>
<td>B</td>
<td>B</td>
<td>B</td>
</tr>
<tr>
<td>11</td>
<td>45</td>
<td>L</td>
<td>B</td>
<td>R</td>
<td>B</td>
</tr>
<tr>
<td>12</td>
<td>29</td>
<td>L</td>
<td>B</td>
<td>B</td>
<td>B</td>
</tr>
</tbody>
</table>

L = left; R = right; B = bilateral. The last column shows correspondence between fMRI results and initial as well as propagated epileptic activity as revealed by the source analysis. The column ‘Spike number’ shows the number of spikes (first spike in the sequence with the EEG baseline of at least 2 s before the spike) that were averaged for the source analysis.
Perisylvian network in CSWS

Our results support previous findings underlining the involvement of the perisylvian region during CSWS. The increased prevalence of CSWS in patients with the perisylvian polymicrogyria (Guerrini et al., 1998) and sporadic association of structural abnormalities in the perisylvian region with Landau–Kleffner syndrome and CSWS (Solomon et al., 1993; De Volder et al., 1994; Tagawa et al., 1999; Huppke et al., 2005) provide clinical evidence to that effect. Consistent with our study, other authors have performed magnetoencephalography/EEG source analysis and have shown that the bilateral spikes generated in or propagate to the perisylvian cortex in many patients with Landau–Kleffner syndrome and CSWS (Morrell et al., 1995; Paetau et al., 1999; Sobel et al., 2000; Paetau, 2009). Positron emission tomography (PET) and single-photon emission computed tomography studies have revealed brain areas of hypermetabolism and hypoperfusion in the perisylvian region and temporoparietal cortex associated with CSWS with and without Landau–Kleffner syndrome (Gaggero et al., 1995; Maquet et al., 1995; De Tiege et al., 2004, 2008; Luat et al., 2005). Morrell et al. (1995) investigated 14 children with Landau–Kleffner syndrome, showing CSWS in all but one. Although bilateral spikes were detected in all cases, the generator of epileptic activity was located unilaterally in the posterior speech cortex (Wernicke’s area, the angular and supramarginal gyri). Patients with this unilateral generator underwent epilepsy surgery in the perisylvian region (multiple subpial transections) with substantial success: 50% of children had recovered age-appropriate speech and 29% improved their neuropsychological abilities, with cessation of CSWS in all cases. Surgical interventions in the perisylvian region have been performed in other studies showing a significant improvement in both EEG and neuropsychological performance (Irwin et al., 2001; Cross and Nevill, 2009; Loddenkemper et al., 2009; Paetau, 2009).

Although the perisylvian region seems to play an important role in the generation of spikes in the clinical tandem of Landau–Kleffner syndrome and CSWS (Solomon et al., 1993; Da Silva et al., 1997; Harbord et al., 1999; Paetau et al., 1999), in patients with CSWS without Landau–Kleffner syndrome the generator may be located in other cortical areas, and the epileptic activity seems to propagate bilaterally to the perisylvian cortex (Gaggero et al., 1995; Maquet et al., 1995; Morrell et al., 1995; De Tiege et al., 2004; Luat et al., 2005). In most cases, the spikes propagate rapidly from one hemisphere to another, so that a focal origin to the spike with secondary bilateral synchrony may be suggested (Kobayashi et al., 1994; Farnarier et al., 1995; Paetau, 2009). Partial motor seizures that originate from different cortical regions (Tassinari et al., 2005), focal EEG activity during both wakefulness and REM sleep (Galanopoulou et al., 2000) and focal cortical areas with hypermetabolism and hypoperfusion individually distributed and well corresponding with the focus of spikes during wakefulness (Gaggero et al., 1995; Maquet et al., 1995) all provide evidence for individual cortical generators in CSWS. In our study, we described individual localization of the initial epileptic activity, which was detected in the perisylvian cortex in only four patients. Epileptic spikes propagated bilaterally to both perisylvian regions supporting the evidence for a secondary bilateral synchrony in CSWS. It is worthwhile to mention that this specific pattern of propagation is identical in symptomatic and cryptogenic cases suggesting that the described bilateral synchrony in the perisylvian region is specific to CSWS rather than any aetiological factor.

The BOLD pattern (Fig. 3) shows an involvement of the prefrontal cortex and the cingulate gyrus. Interpreted in the light of the results of the source analysis, these BOLD responses are likely to represent areas of propagation. This complex pattern of propagation may explain complex neuropsychological deficits that often accompany CSWS. The perisylvian region is involved in acoustic perception and language development (Horwitz and Barun, 2004). The insular cortex and cingulate gyrus are important parts of neuronal networks of working memory, self-control, emotional processing and social cognition (Posner et al., 2007; Rilling et al., 2008; Behrens et al., 2009). Functional abnormalities in these regions may explain the frequent association of CSWS with auditory agnosia, acquired aphasia, aggressiveness, attention-deficit hyperactivity and autistic features (Roulet Perez et al., 1993; Galanopoulou et al., 2000; Deonna and Roulet, 2006; Debiais et al., 2007; Metz-Lutz, 2009). Involvement of the motor cortex in CSWS has been associated with motor impairment in form of dyspraxia, dystonia, ataxia or unilateral deficit (Veggiotti et al., 1999; Galanopoulou et al., 2000) and the appearance of negative myoclonus during wakefulness (Dalla Bernardina et al., 1989). Propagated epileptic activity may disturb function in sensory and cognitive networks in the perisylvian, prefrontal and cingulate cortices and, in such a way, cause neuropsychological deficits (Halasz et al., 2005). Moreover, epileptic spikes may interact with maturational of complex cognitive networks. Perisylvian, prefrontal and cingulate cortices undergo a long developmental process and are especially sensitive to environmental influences and intrinsic physiological perturbations throughout childhood and adolescence (Lenroot and Giedd, 2006; Marsh et al., 2008). Considering consolidation of memory traces during sleep which involve similar cortical structures (Fischer et al., 2005; Takashima et al., 2009), spikes may interfere with the restructuring of cognitive networks in the sensitive phase of development. The reasons for propagation of CSWS to cortical structures that are characterized by a long time frame of intensive synaptic pruning and ongoing progressive myelination of axons, remain to be understood. Our study, however, supports the concept of a link between CSWS and specific developmental abnormalities and residual neuropsychological deficits following successful treatment of this pathological condition (Tassinari et al., 2005).

Involvement of thalamus and striatum in CSWS

A strong association of CSWS and slow sleep provide unequivocal arguments for involvement of the thalamocortical network (Halasz et al., 2005). The majority of children with CSWS either do not have any epileptic activity or show well-localized focal spikes during both wakefulness and REM sleep. However, a pronounced synchronization with bilateral and generalized epileptic activity may be observed in the non-REM sleep (Galanopoulou et al., 2000; Tassinari et al., 2005; Nickels and Wirrell, 2008).
The work of the Steriade group systematically showed that non-REM sleep is characterized by the synchronous bursting-mode of the thalamocortical system, where spikes of the cortical neurons are highly synchronized, driven by the γ-aminobutyric acid (GABA)ergic gating machine of the thalamic reticular nucleus (Steriade, 2003; 2005). Synchronization of epileptiform spikes and slow oscillation during sleep may involve similar mechanisms. It seems likely that, during slow sleep, the cortex is prone to an abnormal synchronizing processes. A close relationship between sleep spindles and spike frequency and transformation of sleep K-complexes into epileptiform paroxysmal depolarization shifts provide an argument for this hypothesis (Gloor et al., 1990; Nobili et al., 1999; Steriade, 2005). In man, primary and secondary generalized paroxysms, as well as bilateral epileptic activity in both generalized and partial epilepsies, have been associated with a strong activation of the thalamus suggesting an involvement of the thalamocortical network in processes of broad pathological synchronization (Aghakhani et al., 2004, 2006; Gotman et al., 2005; Hamandi et al., 2006, Moeller et al., 2008a, b). Moreover, highly synchronous epileptic activity during sleep is commonly accompanied by thalamic activation (Moeller et al., 2008b). We showed that bilateral and generalized epileptic activity in CSWS is commonly associated with a significant activation in the thalamus, as revealed by the group analysis. The prominent role of the thalamus in CSWS has been supported by studies demonstrating that thalamic injuries may cause CSWS (Guzzetta et al., 2009; Kelemen et al., 2006). Taking into account close and rich connections between the thalamus and perisylvian/premotor cortex (Oijermann, 1984; Horwitz and Braun, 2004), it can be assumed that sleep-related increases in pathological thalamic activity may contribute to facilitated synchronization in CSWS and to the described pattern of spike propagation.

Moreover, this study provided evidence for the involvement of the caudate nucleus in CSWS. BOLD decreases in the striatum have been repeatedly found in patients with generalized spike and wave discharges (Gotman et al., 2005; Hamandi et al., 2006; Moeller et al., 2008a, b). The studies on generalized paroxysms have hypothesized that the decrease in the BOLD signal may reflect a reduced corticostriatal drive during epileptic activity, since the caudate nuclei receive strong cortical input from associative areas. Another explanation may be provided by the results from recent studies on rats with idiopathic epileptiform activity which suggested that the basal ganglia act as a remote control system for epileptic spikes. During paroxysms, the cortico-thalamic-palidal network shows rhythmic bursting, whereas striatal output neurons are silenced (Slaght et al., 2004; Paz et al., 2005). It was suggested that the acute drop in firing rate of striatal neurons results from a feed-forward synaptic inhibition, which may contribute to processes of termination of epileptiform activity (Paz et al., 2005, 2007). The decrease in striatal neuronal activity during epileptiform discharges in rats may correspond to the BOLD decreases observed here.

**Default mode network in CSWS**

The first EEG-fMRI study on CSWS was published by De Tiege et al. (2007). The authors investigated a 9-year-old girl suffering from partial seizures and who developed CSWS and neuropsychological deficits. Epileptiform activity was associated with focal activations in the right superior frontal, postcentral and superior temporal cortices as well as deactivations in the lateral and medial frontoparietal cortices, posterior cingulate gyrus and cerebellum, in line with our results. BOLD decreases in the precuneus, retrosplenial cortex, parietal and anterior medial frontal cortices were consistently found in all our patients and in the case investigated by De Tiege et al. (2007). These structures are usually involved in a pattern of deactivation that occurs during the initiation of task-related activity and represent default mode network that is active in the resting brain with a high degree of functional connectivity (Raichle et al., 2001). It has been suggested that the default mode network constitutes a necessary favourable neurometabolic environment for cognitive functions, represents a physiological baseline for processes of attention and working memory and supports dynamical integration of cognitive and emotional processing (Raichle and Mintun, 2006). Abnormal activity in the default mode network and disturbed connectivity between the structures involved may influence task performance and contribute to the pathogenesis of neuropsychiatric disorders such as attention-deficit hyperactivity, Alzheimer’s disease, autism, schizophrenia and depression (Eichele et al., 2008; Broyd et al., 2009). Moreover, altered activity in the default mode network has been associated with fluctuations and disturbance of consciousness (Boly et al., 2008).

It has been suggested that disruption of the resting state activity by pathological processes (e.g. those that give rise to spike) may be related to alterations in cognitive function and this may be the possible mechanism that underlies cognitive deficits in epilepsy (Gotman et al., 2005). Deactivations in the default mode network have been described in awake patients with primary and secondary generalized paroxysms and absence seizures (Aghakhani et al., 2004; Gotman et al., 2005; Hamandi et al., 2006; Laufs et al., 2006; Moeller et al., 2008a, b). These default mode network deactivations may reflect disturbance of awareness or consciousness associated with absence (Laufs et al., 2006; Moeller et al., 2008a). Moreover, in patients suffering from temporal lobe epilepsy, Laufs et al. (2007) found spike associated deactivation of the default mode network, particularly in precuneus and left and right parietal lobes. The authors argue that the transient cognitive impairments and performance deficits observed in temporal lobe epilepsy may be associated with dysfunction of the default mode network.

However, it is important to note that fMRI studies based on correlation analyses such as those mentioned above cannot shed light on the causal relationships between EEG (taken either as purely electrical events or markers of cognitive states) and BOLD patterns. In this regard, models of effective connectivity can be used to study the causal relationships at the neural level from fMRI data such as Dynamic Causal Modelling (Friston et al., 2003). Application of this approach on fMRI acquired during GSW has shown that the state of the precuneus may act as a modulator of the onset (and offset) of the pathological activity (Vaudano et al, 2009). As noted by the authors, this finding relates specifically to the onset of the discharges and does not preclude a reversed causal link between the pathological activity and
deactivation in the default mode network. For example, there could be reinforcement of the default mode network deactivation during sustained discharges, leading to neuropsychological effects.

A prominent pattern of activity in the default mode network has been observed during sleep and has been associated with processes of anatomical connectivity between the frontal and parietal corti cal regions that are necessary for memory consolidation and information processing during sleep (Dang Vu et al., 2008; Horovitz et al., 2009). In our previous study, we demonstrated significant deactivations in the default mode network associated with generalized spike-and-wave discharges in sedated sleeping patients (Moeller et al., 2008b). We suggest that spikes in patients with CSWS may interrupt activity in the default mode network and in such a way intervene with cognitive processes during sleep. Arguments for this suggestion were provided by De Tiege et al. (2008); using longitudinal PET scans acquired before and after successful treatment of CSWS, the authors demonstrated common resolution of default mode network hypometabolism associated with recovery. The authors hypothesized that the neuropsychiologic effects associated with CSWS activity are not restricted to the epileptic focus but spread to connected brain areas via a possible mechanism of surrounding and remote inhibition (Witte and Bruehl, 1999). Our study supports this hypothesis demonstrating spike-associated deactivation in the default mode network independently on the individual focus of epileptic activity.

Limitations of the study

In this study, very frequent epileptic activity was analysed. It could be suggested that the frequent events may violate assumptions of the general linear model and reduce fMRI sensitivity (Jacobs et al., 2008; Horovitz et al., 2009). Nevertheless, accurate source localization using this head model has been demonstrated in different clinical and experimental studies in the past (Lantz et al., 2003a; Michel et al., 2004; Phillips et al., 2005; Sperli et al., 2006). Our study raises issues of interpretation related to the possible effects of sleep-inducing drugs on the relationship between sleep and BOLD, the difficulty of sleep staging and defining a baseline due to the high level of epileptic activity. A number of studies have demonstrated a direct link between sleep depth, level of resting state activity and connectivity in the default mode network and BOLD (Moehring et al., 2008; Horovitz et al., 2009). Furthermore, deactivation in the default mode network may be a product of interaction between epileptic spikes and network providing functional connectivity between anterior and posterior brain regions without any relation to physiological rest (Morcom and Fletcher, 2007). As noted previously the negative BOLD signal changes in the default mode network seem to be non-specific to CSWS (Aghakhani et al., 2004; Gotman et al., 2005; Hamandi et al., 2006; Laufs et al., 2006; Moeller et al., 2008a, b). Whatever the mechanism, the relation between CSWS, memory consolidation and fluctuations of activity in the default mode network has to be investigated in more detail in the future.

Conclusions

Independent of aetiology and individual area of initial epileptic activity, patients with CSWS were characterized by a consistent specific neuronal network of propagation. The activation in the perisylvian/prefrontal network was associated with both activation in the thalamocortical network and deactivation in the default mode network. Since these networks seem prominent in neuropsychological processes and consolidation of memory traces during sleep, a possible influence of epileptic spikes on these networks may explain neuropsychological deficits and developmental abnormalities in CSWS. However, studies are needed that directly investigate information processing during sleep in relation to epileptic activity.

Acknowledgements

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