Mapping Hemodynamic Correlates of Seizures Using FMRI: A Review

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Abstract: Functional magnetic resonance imaging (fMRI) is able to detect changes in blood oxygenation level associated with neuronal activity throughout the brain. For more than a decade, fMRI alone or in combination with simultaneous EEG recording (EEG-fMRI) has been used to investigate the hemodynamic changes associated with interictal and ictal epileptic discharges. This is the first literature review to focus on the various fMRI acquisition and data analysis methods applied to map epileptic seizurerelated hemodynamic changes from the first report of an fMRI scan of a seizure to the present day. Two types of data analysis approaches, based on temporal correlation and data driven, are explained and contrasted. The spatial and temporal relationship between the observed hemodynamic changes using fMRI and other non-invasive and invasive electrophysiological and imaging data is considered. We then describe the role of fMRI in localizing and exploring the networks involved in spontaneous and triggered seizure onset and propagation. We also discuss that fMRI alone and combined with EEG hold great promise in the investigation of seizure-related hemodynamic changes non-invasively in humans. We think that this will lead to significant improvements in our understanding of seizures with important consequences for the treatment of epilepsy. Hum Brain Mapp 34:447-466, 2013. © 2011 Wiley Periodicals, Inc.

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Key words: haemodynamic; seizure; BOLD; EEG-fMRI

INTRODUCTION

Epilepsy affects 5 to 10 persons per thousand [Sander, 2003], and 30% to 40% of patients continue to experience refractory seizures despite medical treatment [Kwan and

bral blood flow, cerebral blood volume, and blood oxygenation resulting in variations in the ratio of diamagnetic oxyhemoglobin and paramagnetic deoxyhemoglobin which can be detected as the blood-oxygen-level-dependent (BOLD) contrast [Kwong et al., 1992; Ogawa et al., 1990]. This intrinsic contrast mechanism provides a means of mapping function-related hemodynamic changes noninvasively over the entire brain with good spatial resolution. In addition to the study of task-modulated effects which are the subject of the majority of fMRI studies, the technique has been used to map brain state-related hemodynamic changes in the resting state [Fox et al., 2009; Gusnard et al., 2001; Laufs et al., 2006] and during sleep [Czisch et al., 2004; Laufs et al., 2007]. EEG was incorporated in the fMRI set up for the first time in 1993 by Ives [Ives et al., 1993], with the later development of

Sander, 2004]. Functional MRI (fMRI) works on the principle that neuronal activity causes regional changes in cere-

Contract grant sponsors: Department of Health's NIHR Biomedical Research Centres funding scheme and The Epilepsy Society; Central and East London NIHR CLRN; and Higher Education Commission of Pakistan (to U.J.C.);

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Received for publication 7 March 2011; Revised 19 July 2011; Accepted 1 August 2011

DOI: 10.1002/hbm.21448

Published online 14 November 2011 in Wiley Online Library (wileyonlinelibrary.com).

EEG-triggered fMRI in epilepsy [Warach et al., 1996]. Simultaneous and continuous recording of good quality EEG and fMRI data in 2001 [Lemieux et al., 2001] further broadened the range of neural events, effects and states that could be studied using fMRI, particularly in the resting state.

The literature on EEG-fMRI has expanded rapidly in the last decade, especially on the study of interictal epileptic discharges (IED) related hemodynamic changes in focal [Archer et al., 2003; Bagshaw et al., 2005; Jacobs et al., 2009; Kobayashi et al., 2006a; LeVan et al., 2010a; Salek-Haddadi et al., 2006] and generalized epilepsy [Hamandi et al., 2006]. This has led to new insights on the effects of epilepsy on neurovascular coupling [Carmichael et al., 2008], localization of epileptogenic zone in focal epilepsy [Federico et al., 2005b; Jacobs et al., 2008; Salek-Haddadi et al., 2006; Thornton et al., 2010a; Zijlmans et al., 2007] and the involvement of epileptic networks for 3 Hz generalized spike wave discharges (GSWDs) [Gotman et al., 2005; Hamandi et al., 2006; Kobayashi et al., 2006b]. However, the delineation of the brain areas producing interictal spikes, known as the ''irritative zone'' (IZ), may not suffice to localize the seizure onset zone (SOZ), which is clinically more important [Luders et al., 2006]. Thus there has been an increasing interest in investigating specific seizurerelated hemodynamic changes using fMRI. Due to methodological and practical issues the data on seizure-related hemodynamic changes is not as extensive as for the interictal state. Our aim is to provide a detailed review of the published literature on mapping seizure-related hemodynamic changes using fMRI alone or combined with EEG. We have described the main methodological constraints, followed by clinical applications, and have discussed the possible future role of the technique in investigating seizures. We will not discuss in detail the technical aspects of EEG recording during fMRI which have been comprehensively discussed in a number of reviews recently [Laufs and Duncan, 2007; Laufs et al., 2008].

For the purpose of this work, we performed a comprehensive search of English literature on Medline Pubmed database with the keywords; ''functional imaging AND epilepsy,'' ''functional imaging AND seizures,'' ''ictal fMRI,'' ''EEG-fMRI,'' ''EEG-fMRI AND epilepsy,'' and ''EEG-fMRI AND seizures'' from 1990 to 2010. In addition we also searched through the bibliographies of the relevant original articles. We selected all original research articles investigating seizure related hemodynamic changes using fMRI alone or combined EEG-fMRI (Table I).

METHODOLOGICAL CONSIDERATIONS FOR MAPPING SEIZURES USING FMRI

Studies using fMRI and/or EEG-fMRI as an investigation tool face a number of methodological issues pertaining to patient recruitment, seizure detection, and data analysis which makes seizure-related fMRI data acquisition complex. Here we will discuss these issues within the specific context of their demonstrated application.

Patient Selection

Patient recruitment for ictal studies using fMRI has been either fortuitous, in cases when seizures have occurred during studies of the interictal state, or targeted. The majority of series published to date have resulted from the former. The targeted selection of patients based on the specific aim of capturing seizures has generally been when seizures were provoked by certain triggers such as fixation-off sensitivity (FOS) [Di Bonaventura et al., 2005; Iannetti et al., 2002; Krakow et al., 2000], music [Morocz et al., 2003], reading [Salek-Haddadi et al., 2009] and photoparoxysmal response (PPR) [Moeller et al., 2009a,b]. Most studies have been single case reports or small series of patients with specific subgroups of epilepsy (Table I). The unpredictable nature of seizures (except when seizures can be triggered) explains this situation.

Vigorous head or body movements inside an MRI scanner may be hazardous from the patient safety perspective [Lemieux et al., 1997] and also adversely affect data quality [Lemieux et al., 2007]. Patient recruitment has, therefore, been limited to specific seizure types which can be scanned without patient safety and data quality concerns.

Seizure Identification Methods

Various means of seizure identification have been applied in fMRI studies including: online and retrospective review of EEG, Electromyography, video recording, observation of ictal semiology (for example: by person inside the scanner room), eye tracking technology, patient response button, and sound/voice recording (Table I). Using simultaneous video-EEG recording is a well established method of seizure identification and video recording can provide additional valuable information to identify seizures when EEG is not helpful [Binnie et al., 1981; Smith, 2005]. The synchronization of EEG and video is pivotal, as there may be a time delay for the electrical activity to propagate from the SOZ to the symptomatogenic zone to produce clinical symptoms [Luders et al., 2006]. Thus depending solely on video or sound recording may be inaccurate or suboptimal. Similarly, if consciousness is affected immediately at the beginning of a seizure, the use of patient signal (e.g. via a button press) may not be a reliable way of identifying seizure onset because the patient may not be able to press the button at all due to impairment of consciousness. Recently a system for simultaneous and synchronized video recording during EEG-fMRI (vEEG-fMRI) has been successfully demonstrated without significant detrimental effects on image quality [Chaudhary et al., 2010]. The combined information from video-EEG can help to identify seizures semiologically and electrophysiologically. Electromyography recordings may also prove helpful during EEG-fMRI studies for atonic/tonic seizures and myoclonic jerks.

Study	Syndrome	Subjects	EEG/ channels	Motion Cut-off (mm)	Seizure activation procedure	Seizure detection	Data analysis approach clinic/results
Adults: fMRI only studies							
[Detre et al., 1995]	FE	$\mathbf{1}$	N		N	OBN	Δ Signal Intensity
[Detre et al., 1996]	FE	$\,1$	N		N	NM	Δ Signal Intensity
[Krakow et al., 2000]	FOS	$\mathbf{1}$	N		$\mathbf F$	ETS	GLM
[Kubota et al., 2000]	FE	$\mathbf{1}$	N		N	OBN	Δ Signal Intensity
[Krings et al., 2000]	FE	$1\,$	N		N	OBN, VR	Δ Signal Intensity
[Morocz et al., 2003]	ME	$\mathbf{1}$	N		M	PRB	GLM, ICA
[Archer et al., 2006]	FE	$\mathbf{1}$	N	$\overline{}$	N	VR	GLM
[Auer et al., 2008]	FE	$\mathbf{1}$	N	0.2	N	OBN	Δ Signal Intensity
[Donaire et al., 2009]	FE	$\mathbf{1}$	N	0.5	N	OBN	Seq.GLM, ICA
Adults: EEG-fMRI studies							
[Iannetti et al., 2002]	FOS	3	Y/32		$\boldsymbol{\mathrm{F}}$	EEG	GLM
[Salek-haddadi et al., 2002]	FE	$\mathbf{1}$	Y/11		N	EEG	GLM
[Salek-haddadi et al., 2003]	JAE	$\mathbf{1}$	Y/nm	$\overline{}$	N	EEG	GLM
[Aghakhani et al., 2004]	IGE	25	Y/21		N	EEG	GLM
[Gotman et al., 2005]	IGE	25	Y/21	$\mathbf{1}$	N	EEG	GLM
[Federico et al., 2005]	$\rm FE$	$\ensuremath{\mathsf{3}}$	a		SD	VR	GLM
[Di Bonaventura et al., 2005]	FOS	$\ensuremath{\mathsf{3}}$	Y/nm		$\boldsymbol{\mathrm{F}}$	EEG	GLM
[Di Bonaventura et al., 2006]	RE	$\mathbf{1}$	Y/nm		N	EEG	GLM
[Di Bonaventura et al., 2006]	FE/IGE	32/11	Y/18	$\overline{}$	N	EEG, VR, PRB	GLM
[Kobayashi et al., 2006]	FE	$\mathbf{1}$	Y/27	$\mathbf{1}$	N	EEG	GLM
[Hamandi et al., 2006]	IGE/SGE	30/16	Y/12		N	EEG	GLM
[Laufs et al., 2006]	JAE	$\mathbf{1}$	Y/29		N	EEG	GLM
[Hamandi et al., 2008]	IGE/SGE	2/2	Y/32		N	EEG	GLM
[Carmichael et al., 2008]	IGE/SGE	2/2	Y/32		N	EEG	GLM
[Salek-haddadi et al., 2008]	ReE	9	Y/11	$\overline{}$	Re	EEG, SR, EMG, PRB	GLM
[Tyvaert et al., 2008]	FE	$\,8\,$	Y/25		N	EEG	GLM
[Donaire et al., 2009]	FE	10	Y/27		N	OBN	Seq.GLM
[Tyvaert et al., 2009]	FE	17	Y/25	$\mathbf{1}$	N	EEG, VR	Seq.GLM
[LeVan et al., 2009]	FE	15	Y/25		N	EEG	ICA
[Chassagnon et al., 2009]	FE	$\mathbf{1}$	Y/27		N	EEG	GLM
[Marrosu et al., 2009]	ME	$\mathbf{1}$	Y/nm	$\overline{}$	N	EEG	GLM
[Thornton et al., 2010]	FE	83	$Y/32 - 64$		N	EEG	GLM, ICA
Children							
[Jackson et al., 1994]	FE	$\mathbf{1}$	N		N		Δ Signal Intensity
[Labate et al., 2005]	IGE	$\mathbf{1}$	Y/18		N	EEG	GLM
[Liu et al., 2008]	EMA	$\overline{4}$	Y/21	$\overline{}$	N	EEG	GLM
[Moeller et al., 2008]	CAE	10	Y/30		N	EEG	GLM
[Moeller et al., 2008]	IGE	10	Y/30		N	EEG	GLM
[Moeller et al., 2009]	IGE/PPR	16/14	Y/30		Ph	EEG	GLM
[Li et al., 2009]	CAE	15	Y/34		N	EEG	GLM
[Moeller et al., 2009]	IGE	$\mathbf{1}$	Y/30	$\overline{}$	Ph	EEG	GLM
[Bai et al., 2010]	CAE	42	Y/21	>3	DR, SD	EEG	GLM
[Moeller et al., 2010]	IGE	12	Y/25		N	EEG	GLM, ICA
[Berman et al., 2010]	CAE	37	$Y/21 - 32$		DR	EEG	GLM
[Moeller et al., 2010]	?IGE	1	Y/32			EEG	GLM
[Moeller et al., 2010]	CAE/JAE	14	$Y/25 - 30$	$\mathbf{1}$	÷.	EEG	Seq.GLM
[Carney et al., 2010]	CAE	11	Y/18		DR	EEG	GLM

TABLE I. Publications on mapping seizure-related BOLD changes using fMRI or EEG-fMRI

^aOne patient had simultaneously recorded EEG.FE = focal epilepsy, FOS = fixation-off sensitivity, ME = musicogenic epilepsy, JAE = juvenile absence epilepsy, IGE = idiopathic generalized epilepsy, RE = Rasmussan's encephalitis, SGE = secondary generalized epilepsy, ReE = reading epilepsy, EMA = eyelid myoclonia with absences, CAE = childhood absence epilepsy, PPR = photoparoxysmal response, N = none, Y = yes, F = fixation, M = music, SD = sleep deprivation, Re = reading, Ph = photic stimulation, DR = drug reduction, OBN = observed by neurologist, ETS = eye tracking system, VR = video recording, PRB = patient response button, SR = sound/voice recording, EMG = electromyography, Δ = change, GLM = general linear model, ICA = independent component analysis, $Seq. = sequential, EEG = electroencephalography.$

Reducing Artifacts in EEG Data Recorded During fMRI

EEG recordings inside an MRI scanner are obscured by two main types of artifacts: gradient artifact (GA) and pulse artifact (PA). For most applications, in particular those relying on the quantitative EEG analysis, it is necessary to remove these artifacts subsequently before any further processing. The GA is produced by the rapidly changing magnetic field used for fMRI acquisition. A number of methods have been devised over the years with varying degrees of efficacy including: frequency domain method [Hoffmann et al., 2000], average image artifact subtraction, average image artifact subtraction and adaptive noise cancellation [Allen et al., 2000], and principal component analysis (PCA) [Negishi et al., 2004; Niazy et al., 2005]. To date the most commonly used method is average image artifact subtraction and adaptive noise cancellation [Allen et al., 2000] which is used both for online and offline EEG artifact removal.

The PA is also often called the ballistocardiographic artifact (BCG). This effect results at least in part from the pulsatile movement of the head and scalp in relation to the cardiac cycle which results in induced voltages (artifacts) in the EEG recording circuit. This effect may make the identification of epileptic events difficult in most patients. Here again methods based on the average artifact subtraction principle [Allen et al., 1998] are the most commonly applied technique for removing PA. Other methods include applying adaptive filtering [Bonmassar et al., 2002], weighted average subtraction [Goldman et al., 2000], median filter template [Ellingson et al., 2004] or statistical algorithms in independent component analysis (ICA) [Srivastava et al., 2005]. All these methods have been validated in a number of studies and provide adequate solutions facilitating the identification of epileptic discharges.

Artifacts in the fMRI Data

The quality of the fMRI data acquired during EEG recording in general and seizures in particular can be severely degraded by a number of factors, namely electromagnetic interference caused by the EEG recording system (passive: distortion from metallic electrodes and leads; active: contamination of MR signal by radio-frequency waves generated by the EEG electronics), head motion, and pulse and respiration related artifacts.

Head motion-related artifact

Head and body movements as small as a fraction of a millimeter can affect fMRI data quality adversely and may make them unusable or impossible to interpret [Hajnal et al., 1994], unless special postprocessing steps are utilized [Friston et al., 1995a]. Motion is integral to most seizure types. Therefore physical means to limit motion such as vacuum cushions can play an important role at the acquisition stage [Benar et al., 2003].

The most common and widely applied motion correction method in fMRI is postacquisition realignment of the fMRI time-series using six rigid body realignment parameters (RP). The realignment parameters are the three rotation angles and three translation shifts that describe the rigid-body relationship between an individual volume in the fMRI time series and the first volume in the series, estimated using a least squares approach and expressed as six time series [Friston et al., 1995a]. Some investigators [Gotman et al., 2005] have used a cut off limit of 1 mm and 1 degree inter-scan motion above which the whole data-set is discarded from the analysis. Each seizurerelated fMRI dataset being considered in this review is potentially unique, in contrast to most cognitive studies in healthy subjects; therefore, another approach is to systematically attempt to extract as much information as possible from every dataset through adapted signal modeling strategies. For example, residual motion-related signal may be treated as confound within the general linear modeling framework. These effects can be modeled based on the motion RPs derived from the scan realignment procedure [Friston et al., 1995a] and by calculating additional linear and nonlinear motion-related effects (using Volterraexpansion of motion RPs) [Hamandi et al., 2006; Salek-Haddadi et al., 2003].

For large motion effects, the inclusion of ''scan nulling'' regressors has also been proposed [Lemieux et al., 2007], instead of discarding the scans, on the basis of the principle that inclusion of nuisance effects into models of the fMRI signal, in addition to the effects of interest, can increase sensitivity [Friston et al., 1995b]. These regressors attempt to explain the signal variation due to severe head motion events (i.e. above a prefixed cut-off value of the inter-scan motion derived from the RP), by representing the effect as a series of four regressors each consisting of a one TR duration Heaviside function corresponding to the scan affected by head motion and the 3 subsequent scans [Lemieux et al., 2007].

Cardiac pulse and respiration related effects: Physiological noise

The cyclical variability of the pulse causes changes in oxygenated blood volume in the brain, and small displacements of the brain tissue from compression and decompression during the cardiac cycle. Similarly, respirationrelated physiological noise also affects fMRI data quality through changes in magnetic field secondary to chest movements and due to changes in $pCO₂$ (partial pressure of $CO₂$ in blood) levels causing fluctuations in BOLD. Though these signal variations are present in all fMRI studies, their effect becomes more significant for EEGfMRI studies of epilepsy in view of the greater imperative of extracting as much information as possible from individual datasets and due to the possible clinical

implications of the result for the patient's management in some cases; this is in contrast with most fMRI studies in which a dataset of suboptimal quality can be replaced with a better one, possibly from a different subject.

Image data correction and regression methods based on the phase of the cardiac cycle relative to the scan acquisition [Glover et al., 2000; Liston et al., 2006] and independent component analysis (ICA) based identification of physiological noise related components and removal [Perlbarg et al., 2007] have been used. For respiration related signals, modeling of respiratory volume per minute and changes in depth and height of respiration as confounds [Birn et al., 2006; van Houdt et al., 2010] have been employed. It has been shown in EEG-fMRI studies of IEDs [Glover et al., 2000; Liston et al., 2006] and alpha rhythms [van Houdt et al., 2010] that inclusion of these additional regressors as confounds improves sensitivity. In addition cardiac effects have also been modeled in fMRI analysis of seizures [Thornton et al., 2010b].

The inclusion of as complete and thorough a model of the effects of motion and physiological confounds such as pulse and respiration on the fMRI signal is the best possible means of avoiding false-positive findings and therefore guaranteeing maximum specificity of the observed BOLD signal changes in relation to neural activity.

fMRI Data Analysis: Seizure-Related BOLD Effects

The overall aim of the analysis is to identify hemodynamic patterns reflecting the epileptic activity. Three types of data analysis approaches have been proposed for this purpose: the identification of regions showing particular patterns of signal fluctuations; the general linear model (GLM); and data-driven such as ICA. Within the GLM framework, the choice and specification of the model embody the investigator's hypotheses. These differ in their underlying assumptions and in particular the degree to which they rely on the use of other data such as synchronously-recorded EEG to reveal these patterns.

Identification of variations in MR signal intensity without reference to EEG or other event marker

Early studies investigating hemodynamic changes during seizures used fMRI without simultaneous EEG or video. In these, fluctuations in signal intensity relative to the mean signal [Detre et al., 1995; Jackson et al., 1994] or the baseline signal [Krings et al., 2000; Kubota et al., 2000] were identified to produce maps of ictal fMRI change. Detre et al. performed a cross-correlation analysis between regions of the earliest signal change including the well characterized focus and the rest of the brain [Detre et al., 1996]. More recently, Auer et al. correlated the signal at every voxel in the brain to an internal reference curve defined by visually examining the signal alterations of the

voxels in the area responsible for seizures to map changes in other regions [Auer et al., 2008].

Detection of variation in signal intensity, as applied in the earlier studies, lacks the concurrent information from EEG on seizure onset and evolution; in addition, mapping of putative ictal hemodynamic changes based on visual identification of regions of raw signal intensity changes may be particularly unreliable due to the effects of motion and MR signal drift.

Correlation analysis: General linear model

The analysis of most fMRI time series data relies on the application of the GLM which is a correlation based technique in a mass univariate (voxel-by-voxel) approach where the signal is expressed as a weighted sum of effects, fitted to the data at each voxel and subsequent mapping of the fitted weight statistics [Friston, 1996]. In practice, it is most commonly implemented in the form of a design matrix encompassing all known effects on the signal either as effects of interest (EOI) or confounds. This is also the case in most EEG-fMRI applications, where the effects associated with epileptic discharges identified on EEG are incorporated into the GLM as a regressor of interest, to map the degree of correlation of this regressor with the measured signal across all voxels. Each EOI to be included as a regressor in a GLM design requires the choice of a mathematical representation, and is convolved with hemodynamic response function (HRF).

A number of questions needs to be considered and explored further in detail to find an appropriate and more realistic way of modeling seizures in fMRI data analysis. For example, can we presume that the hemodynamic changes that take place during a seizure that lasts 10 s correspond to a neuronal event with fixed intensity over that duration? More specifically, can a seizure be realistically represented as an event block analogous to those used in block design cognitive fMRI studies? There is also the issue of the seizure onset and the preictal state; do they correspond to distinct states that fundamentally differ from the baseline or is there a gradual hemodynamic transition from interictal to ictal state? Similarly, can multiple seizures be considered as the same phenomenon and therefore be grouped as a single effect for the purpose of modeling?

Event representation. The EOI are usually represented as box-car functions for example in the fMRI studies following a paradigm such as cognitive studies, or idealized zero-duration stick functions for example in the fMRI studies of random brief events such as IEDs. However, seizures are dynamic in nature, potentially involving a complex sequence of various combinations of brain regions over time. This complexity makes their mathematical representation for GLM model building much more difficult, which is reflected by the variety of approaches used to analyze ictal fMRI data.

In the case of provoked seizures this is somewhat simplified by the possibility of identifying a clear seizure

onset using the timing of provoking factor such as: eye closure in FOS [Di Bonaventura et al., 2006; Iannetti et al., 2002; Krakow et al., 2000], music in musicogenic seizures [Marrosu et al., 2009; Morocz et al., 2003], and reading in reading epilepsy [Salek-Haddadi et al., 2009]. In some case it may be possible to provoke repeated seizures, separated by periods of rest which may therefore be considered repetitions of the same type of event (similar to the block design of conventional cognitive fMRI studies) and represented mathematically as box-car [Iannetti et al., 2002; Krakow et al., 2000; Marrosu et al., 2009; Morocz et al., 2003] or as series of stick functions [Salek-Haddadi et al., 2009]. An important issue which has not been taken into account for provoked seizures is the possible time lag between stimulus presentation and seizure onset, which could allow further exploration of the mechanisms of seizure generation.

In contrast, spontaneous seizures such as focal or absence seizures which occur at random may have uncertainty in the identification of the event onset given the available means. Ictal events must be identified employing different methods (Seizure Identification Methods and Table I), and categorized using the same data. Onset and offset timings of single or multiple seizures or preictal states have been used to define the EOI to be modeled and represented as variable duration blocks (depending upon the seizure duration) to be included in the design matrix as regressors [Salek-Haddadi et al., 2002, 2009; Tyvaert et al., 2008].

Seizures are an evolving phenomena and when seizures are modeled as a single block, though they reveal BOLD changes, the ability to differentiate seizure onset from propagated activity is diminished. Based on this hypothesis, peri-ictal hemodynamic changes have been mapped sequentially; from before the seizure onset (preictal: 120 s before clinical seizure onset [Donaire et al., 2009a], 9 s before the ictal discharge onset on EEG [Tyvaert et al., 2009]) up to prespecified time following the seizure offset (postictal: 60 s after clinical seizure end [Donaire et al., 2009a], 10 s after the EEG discharge offset [Tyvaert et al., 2009]). Donaire et al. divided the preictal, ictal, and postictal blocks into sequential fixed width 10 s blocks each representing a different effect (i.e. independent regressors in the GLM) and compared with a baseline period (20 s temporally unrelated to the seizure). Secondly, the signals in each 10 s block were also compared with that in the contiguous 10 s block within the GLM framework [Donaire et al., 2009a,b]. This approach provides the ability to map and differentiate seizure onset and seizure propagation; however it makes minimum use of the simultaneously recorded EEG. More recently, Thornton and co-workers have modeled seizures by dividing each seizure into three distinct and consecutive phases (representing each as a block of variable durations): early ictal, clinical ictal, and late ictal, based on the onset, duration, and electroclinical evolution over time. The authors argued that this modeling approach makes better use of the available physiological information and allows for the better understanding of

propagation of hemodynamic effects between phases [Thornton et al., 2010b].

Based on the principle that all quantifiable effects should be included in the GLM design matrix to reduce the amount of residual signal variance, IEDs and confounds: RP and pulse regressors have also been represented in the same design matrix as separate regressors and confounds respectively (Head motion-related artifact and Cardiac pulse and respiration related effects: Physiological noise) [Chassagnon et al., 2009; Gotman et al., 2005; Hamandi et al., 2006; Laufs et al., 2006; Salek-Haddadi et al., 2002, 2009; Thornton et al., 2010b; Tyvaert et al., 2008]. As yet, however, there is no consensus on the degree to which such confounds must be modeled.

An important question here is; how should one represent multiple seizures in the design matrix? This is of utmost importance because; unless all seizures recorded are electroclinically stereotyped, they may be manifestations of a variety of underlying epileptogenic networks, in which case they must be represented as different effects (different regressors) in the GLM. Three different approaches have been applied:

- 1. Similar repeated seizures as variable duration blocks in a single regressor [Tyvaert et al., 2008].
- 2. Each epileptic event of variable duration represented as a separate regressor and the event with the longest duration and clear electrical and spatial propagation represented as seizure [Tyvaert et al., 2009].
- 3. Ictal phase-specific effects represented as variable duration blocks across seizures [Thornton et al., 2010b].

Epilepsy-related HRF. Once the events are identified and represented mathematically they must convolve with an informed basis set representing the shape of the $HRF¹$ for the purpose of inclusion in the GLM. The creation of regressors to represent the ictal events themselves (Event representation) and the choice of HRF for ictal events is particularly problematic given that ictal events can last many minutes, and may involve multiple brain regions at different times as demonstrated by seizure semiology and electrophysiology. In addition, we do not know if the BOLD changes associated with ictal events remain

In this work, the use of the conventional ''response function'' terminology (as in HRF, which is derived from cognitive experiments with clear causality) does not imply a causal relationship between epileptic events and putative BOLD signal change. In the context of resting-state EEG-fMRI experiments, the use of features identified on scalp EEG does not guarantee that any fMRI signal change associated with these events will occur after the event. This is due to the possibility of a degree of decoupling due to the limited sensitivity of scalp EEG: for example, one could imagine physiological changes (associated with hemodynamic changes) taking place in the brain before they appear on the scalp and in a time-locked fashion. In the same spirit, we use the "epileptic event related BOLD signal change" terminology in preference to ''BOLD response.''

stationary in a particular brain region once started, or they evolve and diminish before propagating to other brain areas. Thus, longer duration and dynamic nature of seizures increase the difficulty to select a basis set or shape of HRF for seizures.

On the other hand, an important consequence of the univariate aspect of the GLM approach and the choice of basis set is the possibility of accounting for spatial variability in the shape of HRF as it has been shown in interictal and ictal studies that the shape and timing of the peak HRF may deviate from the canonical shape in relation to generalized activity [Bagshaw et al., 2004; Benar et al., 2002; Grouiller et al., 2010; Hawco et al., 2007; Jacobs et al., 2009; Masterton et al., 2010; Moeller et al., 2008a, 2010a; Siniatchkin et al., 2010]. It has also been shown that the shape of HRF may be epilepsy syndrome specific and the use of syndrome-specific HRF may improve the localization of BOLD signal changes [Masterton et al., 2010]. The lesser the flexibility of the basis set the fewer the questions that can be addressed related to the time course of the ictal related changes. For single-term basis sets, such as the canonical HRF, the only question that can be addressed is the sign of the putative BOLD changes. Therefore, interdependence between the temporal and spatial aspects of the analysis is embodied in the necessary choice of HRF for GLM model building, which reflects the investigator's specific hypotheses about the inter-relationship between the ictal phenomenology and the fMRI signals [Archer et al., 2006; Donaire et al., 2009a,b; Hamandi et al., 2006, 2008; Laufs et al., 2006; Salek-Haddadi et al., 2002, 2009; Thornton et al., 2010b].

For more extensive basis sets such as the Fourier set consisting of a linear combination of n sine and n cosine functions, the time course can be evaluated over the specified time window [Salek-Haddadi et al., 2002; Thornton et al., 2010b]. Multiple GLMs each with different gamma functions (time to peak: 3, 5, 7, and 9 s) have also been used to account for some variability in the timing of the HRF [Aghakhani et al., 2004; Chassagnon et al., 2009; Kobayashi et al., 2006b; Tyvaert et al., 2008].

Common features: Mapping group effects. The GLM framework has been used to explore group effects by combining the single-subject analyses using spatial normalization [Penny and Holmes, 2004] for the purpose of making population level inferences [Friston et al., 1999; Worsley et al., 2002]. This is particularly useful to identify shared features and is probably most relevant in stereotypical syndromes such as idiopathic generalized epilepsy (IGE) [Gotman et al., 2005], childhood absence epilepsy (CAE) [Moeller et al., 2008b], and some types of secondary generalized epilepsy (SGE) [Hamandi et al., 2006].

Independent component analysis

ICA is a well-established and powerful data-driven approach based on basic assumptions regarding the properties of the sources responsible for the observed signals, particularly their statistical independence. It can be used to decompose multichannel time series data into spatially or temporally independent components by maximizing their statistical independence. If the signal satisfies the necessary conditions, each independent component represents the activity of one of the sources of the signal [Calhoun et al., 2001; McKeown et al., 2003]. A common assumption in ICA algorithms is that the sources are spatially stationary [McKeown et al., 1998, 1999] which is helpful in separating the signal related sources from noise sources which are presumed to be nonstationary [DeMartino et al., 2007]. However, seizures involve complex spatial and temporal propagation; therefore, the applicability of ICA to such a dynamic process must be questioned in principle. Nonetheless, ICA has been used to study ictal hemodynamic patterns with some interesting results as discussed in section Applications of ICA in Epilepsy, making this assumption of stationarity less relevant.

The components identified by ICA cannot be ranked by degree of importance, in contrast to principal component analysis (PCA). Furthermore for spatial ICA of fMRI time series data, the number of components can be up to the number of scans. Therefore, some form of data dimensionality reduction step is commonly employed either by prespecifying or calculating (using PCA [McKeown et al., 1998]) or using a model of the noise [Beckmann and Smith, 2004] and the number of sources are identified. Furthermore, spatiotemporal characterization of BOLD signals on one hand and fMRI artifacts on the other combined with a pattern recognition technique has also been used to automatically classify components [DeMartino et al., 2007; Rodionov et al., 2007].

Applications of ICA in epilepsy. Morocz et al. [2003] and Donaire et al. [2009b] applied ICA on ictal fMRI time-series data, and visually compared the activity maps with the GLM based statistical maps of BOLD changes. LeVan et al. [2010b] have used an iterative fixed-point method [Hyvarinen and Oja, 2000] to compute independent components (IC), and repeated the decomposition 20 times to identify stable and reproducible sources. They separated ictal components from physiological components, artifacts, residual motion, and cerebral BOLD activity by applying GLM based modeling assumptions whereby: HRF convolved with seizures was fitted to IC time courses in an auto-regressive model and activation maps for ictal components were generated. For multiple seizures in the same patient, single HRF shape was calculated presuming it remains the same for each seizure [LeVan et al., 2010b]. It is also argued that although ICA can reveal BOLD changes and noise separately [McKeown and Sejnowski, 1998; McKeown et al., 1998] it cannot specifically separate the different types of structured noise [LeVan et al., 2010b]. Thornton and colleagues applied cortex based spatial ICA with automated component classification to identify seizure-related ICs spatially concordant with the SOZ [Thornton et al., 2010b].

What are the respective roles of the GLM or ICA-based analyses?

The GLM embodies an approach to fMRI data analysis based on fitting a postulated model of the fMRI time series to the data. It therefore relies entirely on the ability of the investigator to devise a suitable model to allow the identification of the brain regions for which the degree of fit is sufficiently high (in relation to the level of unexplained variance and given the number of voxels at which the degree of fitness is tested). In other words, this approach can only be used to answer questions of the type: in which parts of the brain does the fMRI signal match a prespecified pattern? Assuming the availability of suitable models for each effect, this approach can be used to account for multiple effects simultaneously such as the BOLD changes related to seizure and interictal activity on one hand [Chassagnon et al., 2009; Salek-Haddadi et al., 2009; Thornton et al., 2010b; Tyvaert et al., 2008] and various sources of physiological noise through the inclusion of pulse [Liston et al., 2006], respiration [van Houdt et al., 2010], and motion effects [Salek-Haddadi et al., 2009]. Therefore, this requires the proper identification of events of interest, the mathematical representation of these events in the GLM framework and a choice of hemodynamic kernel as discussed above. Given that the aim of such studies is to reveal BOLD changes specifically related to seizures in any part of the brain and that this depends on the separation between the ictal and interictal states, it is crucial to realize that the GLM approach effectively attempts to explain all fMRI signal variations across the entire brain based on EEG (and possibly video), which is known to have limited sensitivity.

In comparison, ICA of fMRI by not relying on a postulated specific model of the signal time-course, can be seen as a way of freeing the analysis from the bias of scalp EEG. However, this lack of prior model shifts the problem to the interpretation of the often very numerous components. Nonetheless, ICA of fMRI can reveal BOLD patterns that can be associated with seizures through post hoc comparison with EEG [Thornton et al., 2010b]. We envisage that simultaneous intracranial EEG-fMRI will provide important validation data [Vulliemoz et al., 2011].

Validation of the fMRI findings

Given that the aim of fMRI studies of ictal activity is localization, validation requires two elements: a gold standard (intracranial EEG is used as gold standard in most epilepsy surgery programmes) and a method to compare the fMRI maps with the gold standard. Notwithstanding the precise nature of the gold standard, the latter usually requires the application of coregistration of the fMRI maps and gold standard onto a structural MR image of the subject's brain. Because of the complex nature of the fMRI-maps, summarizing the level of spatial agreement can be challenging and various criteria (Table II) have

been applied which can be grouped into the following broad categories:

- 1. Localization of the most statistically significant BOLD change cluster, and also taking into account the localization of other BOLD change clusters [Salek-Haddadi et al., 2006; Thornton et al., 2010b].
- 2. Localization of the cluster corresponding to the earliest BOLD increase, with a cut-off threshold for statistical significance and spatial extent [Donaire et al., 2009a].
- 3. Localization of the most statistically significant BOLD change cluster with a cut-off threshold for spatial extent (greatest number of voxels) [Di Bonaventura et al., 2006].
- 4. Localization of the most statistically significant BOLD change cluster [Tyvaert et al., 2008].
- 5. Localization of all clusters of BOLD change above predefined statistical and spatial threshold [Gotman et al., 2005].
- 6. Localization of the most statistically significant BOLD cluster, and also the most statistically significant cluster in the individual lobes [Hamandi et al., 2006].

In some instances concordance relative to the gold standard is defined (Table II) in terms of Cartesian [Carney et al., 2010; Hamandi et al., 2006; Iannetti et al., 2002; Salek-Haddadi et al., 2003] or cortically informed [Thornton et al., 2010b] distance, while in others it is defined at the lobar level [Salek-Haddadi et al., 2006; Salek-Haddadi et al., 2009; Tyvaert et al., 2009].

CLINICAL APPLICATIONS

EEG-fMRI has been shown to provide new and unique information on the brain networks involved in relation to epileptic seizures. Evidence relevant to assessing the potential clinical value of EEG-fMRI applied to ictal activity, either in relation to localization of the SOZ or syndrome classification, is very limited. To the best of our knowledge, no systematic comparison of the findings of EEG-fMRI with ictal SPECT or PET has been performed to date. Validation in individual cases by comparison with icEEG and or ictal SPECT or PET, is generally encouraging while keeping in mind the limitations of these techniques and of our general understanding of seizure initiation and propagation. No formal assessment of sensitivity and specificity has been done to date, possibly reflecting both the technique's novelty and an early realization of the difficulties in modeling the relationship between EEG and semiology on one hand and fMRI on the other and interpreting the resulting maps. Therefore, we are limited to relatively anecdotal observations. In the published literature the sensitivity in terms of patients who will have a seizure during EEG-fMRI acquisition varies from 10% to 100%, and the sensitivity in terms of patients revealing BOLD changes

^r **M**apping **H**emodynamic **C**orrelates of **S**eizures ^r

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out of patients with seizures during EEG-fMRI acquisition varies from 66% to 100%. This wide-range of sensitivity largely depends on the variable patient selection criteria used in different studies (Table II). Similarly, the ability of EEG-fMRI studies to localize SOZ in focal epilepsy is also quite variable depending upon the various concordance criteria applied (Table II).

Localization of Seizure Onset in Focal Epilepsy

Early case-reports with fMRI alone showed MR signal build up and recovery to baseline in the presumed SOZ (PSOZ) for clinical/subclinical seizures [Detre et al., 1995; Jackson et al., 1994; Kubota et al., 2000]. In another casereport, Krings et al. observed, temporally and spatially distinct, early MR signal change in perilesional cortex before the clinical seizure and an increase in MR signal intensity in eloquent cortex during clinical seizure (symptomatogenic zone) [Krings et al., 2000].

In two case-reports [Kobayashi et al., 2006b; Salek-Haddadi et al., 2002] and a case-series [Donaire et al., 2009a], using EEG-fMRI and GLM-based data analysis approach, time-locked BOLD increase within the PSOZ was revealed during seizures. The amplitude of BOLD signal increase was found to be generally higher in PSOZ in comparison with other areas. In a series of malformations of cortical development [Tyvaert et al., 2008], using GLM based data analysis approach, the spatial localization of seizurerelated hemodynamic response was shown to be different in various pathological lesions. In nodular heterotopia (four of four cases) the maximum BOLD increase was observed in the overlying cortex of heterotopia whereas in focal cortical dysplasia (FCD) (two of two cases) and band heterotopia (two of two cases) maximum BOLD increase was found within the structural lesion, however, other smaller and less significant clusters of BOLD signal change were also observed remotely [Tyvaert et al., 2008]. Thornton et al. [2010b] investigated seizures by dividing into three phases (see Event representation section for details). Using GLM based data analysis approach they showed significant BOLD changes in seven of nine cases, and the concordance of BOLD localization with the SOZ (defined using icEEG) was found to be greatest for the early ictal phase and the combined phases (see Fig. 1). Moreover, they argued that as the ictal activity spreads on EEG, the

BOLD changes should also evolve temporally. Therefore, BOLD changes during clinical ictal and late ictal phases discordant with SOZ may represent propagated activity. Using ICA on the data from the same group of patients, it was demonstrated that ICs were spatially concordant with SOZ in all cases (confirmed in eight of nine cases with icEEG), making an argument that ICA is helpful to reveal ictal BOLD changes when GLM based approach fails [Thornton et al., 2010b].

It has also been shown that the spatial extent of BOLD signal change was larger for the combined effect of multiple seizures than single seizures [Kobayashi et al., 2006b], and the BOLD changes for seizures were widespread in comparison to the BOLD changes for IEDs [Kobayashi et al., 2006b; Tyvaert et al., 2008]. The significance of BOLD signal decrease is not clearly known, however, additional areas of BOLD signal decrease during seizures were observed; within perilesional cortex and opposite hemisphere of brain [Kobayashi et al., 2006b]; within PSOZ [Donaire et al., 2009a]; within surrounding areas of BOLD signal increase and remotely from the PSOZ/SOZ [Thornton et al., 2010b; Tyvaert et al., 2008]. These remote areas were often part of the so-called default mode network [Raichle et al., 2001], in common with many studies of IEDs [Laufs et al., 2006; Salek-Haddadi et al., 2006].

Preictal Changes, Seizure Evolution, and Propagation in Focal Epilepsy

Federico et al. focused specifically on preictal BOLD changes in three cases and revealed BOLD increase (two of three cases) and decrease (one of three cases) starting before the clinical seizure, however, preictal BOLD increase $(\sim4 \text{ min})$ was concordant with SOZ in only one case [Federico et al., 2005a]. Using series of sequential fixed-width block model for seizure representation in the GLM framework, Donaire et al. [2009a] described widespread BOLD decrease 10 to 26 s before the first BOLD increase which in itself preceded the first clinical (6–52 s) or electrographic change (8–12 s) in 5 of 10 cases. In contrast, using a similar model to Donaire et al.'s, Tyvaert et al. [2009] found that on average BOLD changes started 5.2 ± 2.6 s after EEG-onset of seizure and returned to baseline 28.8 \pm 12.9 s after EEG onset in 10/17 cases, roughly in line with what would be expected in response

Figure 1.

Hemodynamic changes revealed by EEG-fMRI for a single seizure in a patient with FCD in left superior frontal gyrus. (**a**) BOLD clusters overlaid on T1-weighted MRI fused with a CT-scan containing subdural grids, (canonical GLM analysis). (**b**) BOLD clusters overlaid on T1-weighted MRI fused with a CT-scan containing subdural grids, (fourier basis set GLM analysis). (**c**) BOLD clusters overlaid on a surface rendering of T1-weighted MRI with subdural grids, (canonical GLM based approach). (**d**)

Time-course of the BOLD signal change over the course of the seizure. (**e**) ICA clusters overlaid on T1-weighted MRI fused with a CT containing subdural grids. (**f**) ICA component illustrated in (e) overlaid on a surface rendering of the T1-weighted MRI. (**g**) Time-course of the component illustrated in e) over the whole recording with the seizure indicated by the red box [Thornton et al., 2010b]. [Color figure can be viewed in the online issue, which is available at wileyonlinelibrary.com.]

to a physiological block stimulus. The initial area of BOLD increase was found to be concordant with the SOZ/PSOZ (Table II) [Donaire et al., 2009a; Tyvaert et al., 2009]. Moreover, by relaxing the statistical threshold additional BOLD change clusters concordant with SOZ/PSOZ were noticed in 2 of 17 cases before ictal onset on EEG. [Tyvaert et al., 2009]. They argued that other areas of BOLD signal increase demonstrated propagation of seizures as it spreads in time. Though these observations were made in small studies, this may point towards EEG-fMRI's ability to identify preictal hemodynamic changes even when epileptic discharges are not yet visible on scalp EEG.

Seizure-Related Networks in Idiopathic Generalized Epilepsy

Salek-Haddadi et al. [2003], using GLM based data analysis approach, described hemodynamic involvement of deeper structures of the brain during absence seizures, identifying two patterns of simultaneous BOLD change; BOLD increase $(+3)$ % from baseline) in thalamic nuclei and symmetrical, widespread decreases in cerebrum, maximum in the frontal lobes $(-8\%$ from baseline) in relation to four long (mean duration: 30 s) absence seizures. Subsequent group studies of IGE patients [Aghakhani et al., 2004; Gotman et al., 2005; Hamandi et al., 2006; Laufs et al., 2006] also revealed common patterns of BOLD decrease in the frontal and parietal cortices, posterior cingulate, caudate and precuneus, and predominant BOLD increase in the thalamus, mesial frontal region, and insulae (see Fig. 2).

The areas involved in GSWD-related cortical BOLD decrease resemble the default mode network [Raichle et al., 2001]. The BOLD signal decrease in these areas, in relation to GSWDs, can be argued to represent the suspension of physiologic conscious rest or an fMRI signature of the negative clinical phenomenology of absences during which cognitive processes are impaired [Laufs et al., 2006]. The studies of absence seizures in animal models [Danober et al., 1998] and in humans using transcranial Doppler, H2⁽¹⁵⁾O and PET [Diehl et al., 1998; Prevett et al., 1995] and invasive recordings have also pointed towards the involvement of corticothalamic network in the generation of absence seizures [Avoli and Gloor, 1981, 1982a,b; Avoli and Kostopoulos, 1982; Avoli et al., 2001]. However, EEG-fMRI noninvasively provides the evidence for the association of the cortico-subcortical/striatocorticothalamic network with GSWD.

The effect of duration of GSWDs on the spatial distribution of BOLD changes is complex. Aghakhani et al. showed that the BOLD changes for shorter $(<3 s)$ and longer (>3 s) bursts of GSWDs were similar in three of four cases and one case did not reveal any BOLD change for shorter discharges [2004]. On the contrary, it has also been discussed that BOLD changes were of greater magnitude [Carney et al., 2010] or tended to be more frequent in the thalamus [Hamandi et al., 2006; Li et al., 2009] when GSWDs were of longer duration and higher in number. This may be due to improved signal to noise ratio or may reflect the role of thalamus in maintaining the GSWD [Avoli et al., 2001] for longer duration discharges.

Studies using simultaneous EEG-fMRI and perfusion mapping techniques (arterial spin labeling) have shown a positive correlation between BOLD changes and cerebral blood flow irrespective of the sign of BOLD change. This association is stronger during GSWDs than during background activity suggesting preserved (i.e. within normal limits) neurovascular coupling [Carmichael et al., 2008; Hamandi et al., 2008].

Timing of the BOLD changes relative to the EEG onset of GSWD is also under-debate. Using conventional GLM based data analysis approach: GSWDs modeled as box-car and convolved with a standard HRF [Glover et al., 2000], it was shown that BOLD increase in the thalamus, occipital cortex, cerebellum, temporal lobes, insula, caudate, and pons; and decrease in the parietal cortex, precuneus, cingulate gyrus, basal ganglia, and pons started after EEG onset, and BOLD increases peaked earlier than BOLD decreases [Aghakhani et al., 2004; Bai et al., 2010; Berman et al., 2010; Carney et al., 2010; Hamandi et al., 2006; Moeller et al., 2008b]. However, it is argued that this analysis approach may not be perfect physiologically; assuming electrophysiological and hemodynamic changes during seizures start and end abruptly. Therefore, by modeling the HRF before the EEG onset of GSWD in the GLM framework [Moeller et al., 2008a], and by investigating the mean time course of GSWD related hemodynamic change in specific areas (selected on the basis of location of significant BOLD cluster) [Bai et al., 2010; Carney et al., 2010], it was shown that temporal pattern of BOLD changes was different from the conventionally reported pattern. The BOLD increase in the thalamus, [Moeller et al., 2008a], parietal cortex and precuneus [Carney et al., 2010] preceded the GSWD onset on EEG. Bai and colleagues also suggested that the BOLD increase in orbital frontal cortex, frontal polar, cingulate, parietal cortex, precuneus, and occipital cortex preceded the GSWD onset on EEG followed by BOLD decrease in the same areas lasting 20 s after GSWD [Bai et al., 2010]. Animal models of absence seizures investigated with optical imaging also showed changes in concentration of oxy/deoxy-hemoglobin before the onset of GSWD [Roche-Labarbe et al., 2010], illustrating all the more need to correlate the findings of animal models with human studies. The inconsistency of temporal evolution of GSWD-related BOLD changes in IGE may result from variable modeling strategies applied [Bai et al., 2010; Moeller et al., 2008a], different types of epileptic discharges (polyspikes and waves and GSWDs) and agerelated biological differences between adults [Hamandi et al., 2006] and children [Moeller et al., 2008a]. The pre-GSWD BOLD changes may reflect early electrophysiological changes which are not evident on scalp EEG, however, this needs to be investigated further.

Moeller et al. investigated the patient specific BOLD changes for absence seizure, using series of sequential

Figure 2.

BOLD clusters correlated with 3 Hz GSWDs in IGE patients revealed by group analysis. Red cluster reveal activations and green clusters reveal deactivations [Hamandi et al., 2006]. [Color figure can be viewed in the online issue, which is available at wileyonlinelibrary.com.]

fixed width blocks to represent seizures in the GLM framework, and found that BOLD changes were consistent for several absences within one patient but varied across patients [Moeller et al., 2010a]. Moreover, the BOLD changes in various anatomical areas; thalamus, cortex and caudate, were dynamic (evolving at different time points) throughout the duration of absence seizure rather than being static [Moeller et al., 2010a]. They also showed that the BOLD increase in cortex (six of nine cases) and decrease in caudate (five of nine cases) preceded the BOLD increase in the thalamus (nine of nine cases) [Moeller et al., 2010a]. These cortical, absence seizure-related, focal BOLD increases [Bai et al., 2010; Carney et al., 2010; Moeller et al., 2010a] are consistent with the cortical focus theory of initiation of absences [Meeren et al., 2002; Polack et al., 2007; Vaudano et al., 2009; Westmijse et al., 2009].

Seizure-Related Networks in Other Forms of Epilepsy

Seizure-related networks explored in fixation-off sensitivity using EEG-fMRI have revealed BOLD increase in parieto-occipital regions correlated with 2.5 to 3 Hz activity triggered by eye-closure [Di Bonaventura et al., 2005; Iannetti et al., 2002; Krakow et al., 2000]. In two case reports on musicogenic epilepsy, it was shown that BOLD networks involved PSOZ: right gyrus rectus and left anterior temporal lobe [Morocz et al., 2003], and right dorsal frontal cortex and right temporal lobe [Marrosu et al., 2009] during presentation of epileptogenic music. Morocz et al. argued that the BOLD changes in the left anterior temporal lobe represented the seizure focus (confirmed by ictal-EEG and ictal-SPECT) and that the BOLD changes in the ventral frontal lobe and right gyrus rectus were related to presentation and listening of epileptogenic music, in line with previous PET studies [Morocz et al., 2003].

For pseudo-absences (three cases) in frontal lobe epilepsy, focal BOLD increase was revealed in the mesial frontal cortex correlated with GSWDs which was not seen in typical absences (six cases) in the same study [Di Bonaventura et al., 2006]. However, more recent studies on absences in children have also shown focal BOLD increases in the cortex (Seizure-Related Networks in Idiopathic Generalized Epilepsy), therefore, it needs to be further investigated as to whether these different hemodynamic changes between typical and pseudo-absences can be applied as a differentiating point. Similarly, comparison of the GSWD-related BOLD patterns in patients with primary and secondarily generalized epilepsies revealed many similarities in the distribution but greater inter-individual variability in cases of secondary generalization, and did not lead to reliable classification of individual cases [Hamandi et al., 2006].

In patients with reading epilepsy, BOLD networks involving the motor and premotor cortex, striatum, mesial temporal lobe, and thalamus were revealed in correlation with orofacial reflex myoclonus and EEG discharges triggered by reading. These changes were also differentiated from the BOLD activations observed during language and facial motor tasks [Salek-Haddadi et al., 2009]. EEG-fMRI has also helped to demonstrate the coexistence of separate epileptic networks for GSWDs and for focal facial twitching in a single patient, and the BOLD changes for the latter helped to ascertain that the FCD previously seen on MRI was epileptogenic [Chassagnon et al., 2009].

In children with photoparoxysmal response (PPR) (six cases; $4 = IGE/CAE$, $2 = PPR$ only) [Moeller et al., 2009a], focal BOLD increases were observed in premotor and parietal cortex (adjacent to intraparietal sulcus) 3 s before PPR (corresponding to gamma activity on EEG), followed by BOLD decrease in the same areas during PPR. It is possible that the early BOLD changes in parietal cortex indicate its possible role in the generation of PPR, or may reflect it being part of the frontoparietal visual networks responsible for

saccades and visual attention, which needs to be elucidated further [Moeller et al., 2009a]. Additionally, most of the subjects (five cases) did not show any BOLD changes in the thalamus [Moeller et al., 2009a], and it can be argued that the thalamus may play a less important role in the generation of PPR in comparison to typical GSWD in IGE for which an intact thalamocortical network is deemed necessary [Avoli et al., 2001].

Liu and colleagues, investigating children with eyelid myoclonia associated with absences (EMA) showed BOLD increase in thalamus, temporal lobes, midline structures and BOLD decrease in parietal and frontal cortex. This epileptic network in EMA is similar to the epileptic network identified in IGE, which can be explained by the presence of GSWDs both in EMA and IGE [Liu et al., 2008].

Hemodynamic Mapping of Cognitive Networks During GSWDs

The hemodynamic changes for GSWDs associated with cognitive impairment have been investigated only recently using various attention tasks during EEG-fMRI [Bai et al., 2010; Berman et al., 2010; Moeller et al., 2010b]. BOLD increases in thalamus, frontal cortex, primary visual, auditory, somatosensory, and motor cortex, and BOLD decreases in the lateral and medial parietal cortex, cingulate gyrus, and basal ganglia have been revealed for GSWDs during which behavioral performance was poor (four of nine cases, 100% error rate). In comparison, for GSWDs during which behavioral performance was good (two of nine cases, 0% error rate), no significant GSWDrelated BOLD changes were observed [Berman et al., 2010]. These findings provide some evidence that activity in structures thought to be important during attention tasks [Riccio et al., 2002] can be affected during GSWDs. The impairment of consciousness during GSWDs and the associated BOLD changes is a complex question which needs to be explored systematically in the future.

The Role of EEG-fMRI on Seizures in Diagnosis and Management of Epilepsy

In single case reports, EEG-fMRI has also demonstrated the ability to help in diagnosis and management of epilepsy. Spatial location of BOLD changes associated with myoclonic jerks of right foot helped to localize an FCD from left frontal lobe which was confirmed with intraoperative cortical mapping as SOZ and resected resulting in complete abolition of seizures [Archer et al., 2006]. In another patient with refractory epilepsy, who had undergone epilepsy surgery twice without any appreciable success, fMRI showed widespread BOLD changes involving the cortex, caudate nucleus, thalamus and other areas during the seizure (a pattern known to be present in generalized epilepsy). Consequently, the antiepileptic drugs were changed to control generalized epilepsy and seizure fre-

quency reduced from 10/day to 1/month [Auer et al., 2008].

PERSPECTIVES AND CONCLUSIONS

What is the future role of EEG-fMRI? Given that seizures happen randomly with likely head or body motion, it would seem that the application of ictal EEG-fMRI will be limited to a subset of cases with very carefully defined selection criteria. However, we believe that such studies will provide crucial new insights into the spontaneous transition from the interictal to ictal state in humans, complementing invasive EEG studies with their limited spatial coverage.

The most suitable candidates for such investigations are patients with daily seizures and with seizure types: absence seizures, myoclonic jerks localized to limbs, simple motor/sensory seizures, subclinical/electrographic seizures and complex partial seizures without large head movements (Table II). To identify seizures, synchronized video-EEG inside the MRI scanner along with simultaneous testing for awareness by performing cognitive tasks would provide the most valuable data.

Methodologically robust and consistent strategy which can separate seizure-related BOLD changes from artifactual changes need to be applied to analyze seizure-related fMRI data. Considering the clinical importance of EEG to identify seizures, we suggest the use of EEG-based GLM, using a system of event representation of seizures as dynamic entities [Thornton et al., 2010b], as a first line approach. Further exploration could proceed using a more flexible model, such as Fourier expansion [Thornton et al., 2010b] or series of short, uniform blocks spanning the seizure or beyond [Donaire et al., 2009a]. The value of purely data-driven approaches such as ICA for the identification of ictal BOLD patterns will probably require a much better characterization of ictal hemodynamic patterns.

EEG-fMRI studies in focal epilepsy have demonstrated the technique's ability to localize the SOZ confirmed by icEEG [Thornton et al., 2010b]. The future potential of EEG-fMRI to provide a noninvasive way of better planning the placement of subdural grids and intracranial electrodes to outline SOZ has also been proposed [Thornton et al., 2010b]. However, the clinical value in the context of technique's sensitivity and specificity remains to be assessed prospectively which can be very complex considering the way multiple test are done [Knowlton, 2006]. EEG-fMRI studies on GSWDs have progressed more rapidly, possibly due to the presumed more uniform underlying phenomenology. The identification of preictal BOLD changes and BOLD networks for various seizure types using EEG-fMRI has improved our knowledge of the underlying epileptic networks such as cortico-thalamic network in IGE. However, the effects of seizures on the networks involved in cognition and maintaining awareness remains to be explored properly.

Future studies exploiting fMRIs capability to map activity patterns over the entire brain at the second time scale may improve our understanding of seizure initiation and propagation and help us answer clinically relevant questions.

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