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*CORRESPONDENCE Federico Giove, Implications federico.giove@uniroma1.it

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Towards high-resolution quantitative assessment of vascular dysfunction

Maria Guidi¹, Giovanni Giulietti², Emma Biondetti^{3,4}, Richard Wise^{3,4} and Federico Giove^{1.5}*

¹Museo Storico della Fisica e Centro Studi e Ricerche Enrico Fermi, Rome, Italy, ²Neuroimaging Laboratory, Fondazione Santa Lucia IRCCS, Rome, Italy, ³Institute for Advanced Biomedical Technologies, University "G. d'Annunzio" of Chieti-Pescara, Chieti, Italy, ⁴Department of Neuroscience, Imaging and Clinical Sciences, University "G. d'Annunzio" of Chieti-Pescara, Chieti, Italy, ⁵Laboratory of Neurophysics and Neuroimaging (NaN), Fondazione Santa Lucia IRCCS, Rome, Italy

Neurovascular alterations are increasingly recognized as a key feature of many brain diseases. They can manifest as a reduction in resting cerebral blood flow or cerebrovascular reactivity (CVR) in the whole brain or in specific regions, depending on the underlying condition. Neurovascular impairment is observed in hypertension, Alzheimer's disease, stroke, multiple sclerosis and cerebral small vessel disease. Magnetic resonance imaging (MRI)-derived CVR mapping is a reliable marker of vascular dysfunction and has been performed mainly at standard functional MRI (fMRI) resolutions of 2-3 mm using the blood oxygen level dependent (BOLD) contrast. However, vascular alterations may occur at a finer scale (i.e., in the capillary bed) which would be better characterized with smaller voxel sizes. Capillaries in gray matter deliver oxygen and glucose to neural tissue and are arranged in a mesh structure, with variable density across the cortical depth. Given that the human cortex is, on average, 2.5 mm thick, submillimetric voxel sizes are effective in increasing the spatial specificity of measurements of hemodynamic and metabolic changes. Novel MRI sequences offer the possibility to map physiological parameters at high resolution with relatively simple experimental setups. In particular, pairing the BOLD acquisition with a contrast sensitive to blood volume changes, while administering a mild hypercapnic challenge, allows for simultaneous mapping of CVR, cerebral metabolic rate of oxygen consumption and other relevant parameters at a high resolution and can be performed at the clinical field strength of 3 T. We propose that this approach will help provide crucial insights into vascular impairment.

KEYWORDS

BOLD, fMRI, VASO, calibrated fMRI, CMRO₂, CVR, SVD, AD

Abbreviations: AD, Alzheimer's disease; ASL, arterial spin labeling; BBB, blood-brain barrier; BOLD, blood oxygen level dependent; CAA, cerebral amyloid angiopathy; CBF, cerebral blood flow; CBV, cerebral blood volume; CMRO₂, cerebral metabolic rate of oxygen consumption; CO₂, carbon dioxide; CSF, cerebrospinal fluid; CVR, cerebroxacular reactivity; EtCO₂, end-tidal CO₂; fMRI, functional magnetic resonance imaging; GM, gray matter; GRE, gradient echo; M, calibration parameter; MCI, mild cognitive impairment; MRI, magnetic resonance imaging; NVU, neurovascular uncoupling; PaCO₂, arterial partial pressure of CO₂; SE, spin echo; SNR, signal-to-noise ratio; SS-SI-VASO, slice-saturation slab-inversion VASO; SVD, small vessel disease; T1, longitudinal relaxation time; TE, echo time; TR, repetition time; VASO, vascular-space occupancy; WM, white matter.

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1 Introduction

Neurodegenerative diseases represent a huge socio-economic burden in aging societies. Magnetic resonance imaging (MRI) offers information about different brain features non-invasively and it is widely employed in the detection of structural damage. Functional alterations, however, may initiate years before anatomical alterations and are important in the picture of disease pathophysiology. Therefore, reliably mapping vascular dysfunction and identifying early biomarkers should aid prevention and catalyze the development of therapies.

MRI contrast based on the blood oxygen level dependent (BOLD) effect is widely exploited in brain research. BOLD-based functional MRI (fMRI) is largely confined to clinical or basic research due to its nonquantitative nature and its dependency on multiple parameters, which are related partly to baseline tissue structure and physiology [1] and partly to underlying dynamic changes in cerebral blood flow (CBF), cerebral blood volume (CBV) and cerebral metabolic rate of oxygen consumption (CMRO₂). Other fMRI techniques that have found a comparatively large application in clinical studies are arterial spin labeling (ASL), for quantifying perfusion, and cerebrovascular reactivity (CVR) mapping based on BOLD or ASL. These have been applied to the study of aging, stroke, tumor, dementia, multiple sclerosis, brain injury, arterial stenosis, and more [2].

Other quantitative, less explored, fMRI techniques include vascularspace occupancy (VASO) [3], able to quantify changes in CBV, and calibrated fMRI, which relies on a simple model of the BOLD signal for extracting CMRO₂ [4-6]. These techniques have found limited application in disease states, for reasons including the lack of standardized protocols, limited contrast-to-noise and more difficulties in data acquisition and processing compared to BOLD. In this perspective article, we propose that VASO and calibrated fMRI at high resolutions can add valuable information to the picture of vascular impairment because of their straightforward physiological interpretation and their closer link to the microvasculature compared to BOLD [7]. Indeed, many neurodegenerative diseases are accompanied by (or originate from) a microvascular dysfunction, which can be assessed by observing the effects on microvascular parameters (CBF and CBV) or metabolic parameters (CMRO₂), as reported, for example, by Klinkmueller et al. in the case of Huntington's disease [8].

For microvascular assessment, imaging at submillimetric resolution is needed, since the average human cortical thickness is about 2.5 mm [9] and, thus, gray matter (GM) voxels having similar or larger sizes are typically affected by severe partial volume contamination by the surrounding cerebrospinal fluid (CSF) and white matter (WM), which has a different vascularization. Moreover, in terms of vascular density and function, the cortex is organized in a layered structure [10] and it has been shown that voxel sizes of about 0.8 mm are capable of distinguishing layer-specific activation in certain areas based on haemodynamic responses [11].

Calibrated fMRI is currently limited in resolution. In order to calibrate the BOLD signal, in addition to the BOLD contrast, a "vascular" contrast needs to be acquired (usually ASL for mapping CBF) while modulating the arterial partial pressure of carbon dioxide (PaCO₂). The experiments are typically performed at low image resolutions (e.g., $3.5 \times 3.5 \times 3-8$ mm³) [12], which are often dictated by the low signal-to-noise ratio (SNR) of ASL sequences.

VASO-fMRI, on the other hand, has been recently proven to be feasible at submillimetric resolutions (0.8 mm) at the clinical field strength of 3 T [13], therefore it represents a good candidate for microvascular state assessment in disease. Additionally, in the framework of calibrated fMRI, it has been shown that the term related to CBF changes can be replaced by a term related to CBV changes (obtainable with VASO) via Grubb's law, while assuming different coupling exponents for venous and total CBV changes [14, 15]. Therefore, VASO can be used to obtain submillimetric maps of CMRO₂ changes [14].

This perspective article is structured as follows: Section 2 focuses on disease states characterized by microvascular impairment; Section 3 describes the parameters that can be derived from MRI in order to assess microvascular dysfunction in such diseases; Section 4 proposes an acquisition protocol and functional paradigm for mapping the parameters listed.

2 The microvascular component in neurodegenerative diseases

While submillimetric VASO and calibrated fMRI have been proven feasible for the characterization of the microvascular component in healthy individuals [13, 14], we suggest that the same methodology may be useful in the study of disease states. A dysfunctional microvasculature is indeed common in many pathological conditions, such as small vessel disease (SVD), vascular dementia, Alzheimer's disease (AD), ischemic stroke, brain tumors, cerebral amyloid angiopathy (CAA), mild cognitive impairment (MCI), Huntington's disease, Parkinson's disease, among others [16]. We focus here on SVD, AD and brain tumors as case studies for investigating microvascular impairment.

2.1 Small vessel disease

Cerebral small vessel disease includes a wide spectrum of cerebrovascular diseases that involve endothelial dysfunction of capillaries, small arteries and small veins in the brain, leading to blood-brain barrier (BBB) dysfunction, impaired vasodilation, vessel stiffening, dysfunctional blood flow and interstitial fluid drainage, WM rarefaction, ischemia, inflammation, myelin damage, and secondary neurodegeneration [17]. The underlying neurobiological mechanisms are, to date, not clear, and SVD is generally assessed by looking at macroscopic or mesoscopic such as WM hyperintensities, manifestations. lacunas. microbleeds, perivascular spaces, or small subcortical infarcts. In terms of hemodynamic measures, a reduction in CVR and CBF has been observed [18]. Given the microvascular nature of the disease, mapping microvascular reactivity and the possible related metabolic impairment can help clarify the underlying mechanisms and pathological effects of the observed alterations [19].

2.2 Alzheimer's disease

AD is characterized by amyloid- β accumulation and impaired CVR [20], but the relationship between these alterations is not clear.

One hypothesis is that amyloid- β accumulation may directly induce vascular dysfunction by impairing vasorelaxation, but therapies aimed at amyloid- β plaque reduction are not effective in reversing cognitive decline [21].

Another hypothesis identifies the microvasculature and related inflammatory processes as fundamental in the pathogenesis of AD [22]. In particular, a CBF reduction is currently the earliest known change associated with the disease, and the mechanism behind it might be a pericyte-driven constriction of capillaries [23]. It is hypothesized that this constriction and the associated hypoperfusion are the precursors of amyloid- β accumulation via upregulation of the BACE1 enzyme. Capillary dysfunction and microvascular CVR reduction were additionally observed to be associated with symptom severity [24, 25], and resting CMRO₂ was found to be reduced in a calibrated BOLD study in AD patients [26].

2.3 Brain tumors

Brain tumors are accompanied by strong alterations in the vasculature (e.g., angiogenesis), which often lead to a local disruption in the neurovascular coupling, also termed neurovascular uncoupling (NVU). Even low grade brain tumors show some degree of NVU [27]. In practice, this may cause a reduced or absent BOLD response to functional tasks in the region affected, even if neuronal activation is present [28]. Distinguishing regions of NVU from regions effectively lacking neuronal response is very important in presurgical planning to avoid excess surgical resection [27, 28]. In place of task-based BOLD fMRI, CVR mapping is generally preferred as it relies on a purely vascular stimulus. CVR mapping was found to be superior (i.e., able to identify NVU regions also in low and intermediate brain gliomas) to perfusion mapping with T2* dynamic susceptibility contrast [27]. Increasing spatial resolution in CVR maps and adding a vascular contrast (such as CBVweighted) is important for precise localization and characterization of the tissue affected.

3 Quantitative parameters for high resolution mapping of vascular dysfunction

In the following, we list quantitative fMRI-derived parameters useful in the study of neurovascular disease and that can be mapped at high resolution at 3 T.

3.1 Cerebral blood volume

Mapping of CBV with MRI without the use of contrast agents can be achieved with vascular-space occupancy [3]. VASO is an inversion recovery technique exploiting the different T_1 of blood and tissue ($T_{1,tissue} < T_{1,blood}$) for creating a contrast sensitive to CBV changes. In the original VASO formulation the images were acquired at the time of blood nulling [3], while more recent implementations showed that different inversion times can be used, as long as a difference in the T_1 weighting of the two compartments is achieved [29, 30]. When the vasculature dilates, either following a metabolic demand of the tissue or a vasodilatory stimulus, the volume of blood in a responding voxel increases. VASO measures quantitative changes in blood volume, which can be expressed as a percentage change or in volume units, but VASO does not quantify baseline CBV.

The slice-saturation slab-inversion VASO (SS-SI-VASO) [31], developed at 7 T, permits the acquisition of gradient-recalled echo (GRE)-BOLD and VASO contrasts interleaved by adding a second excitation pulse at each repetition time (TR). VASO demonstrated a high specificity for the capillaries, arterioles and intracortical arteries [32, 33], making it a valuable tool for high-resolution fMRI [34]. Recently, the VASO implementation acquiring BOLD volumes concomitantly has been adapted for high-resolution applications at 3 T, where it has been shown capable of distinguishing layerdependent activation [13], thus reflecting a microvascular sensitivity.

Using the VASO sequence requires some effort in setting up a working protocol, dealing with the lower SNR and longer TRs compared to BOLD, as well as a less standardized data processing. Examples of protocols for submillimetric acquisitions can be found at https://github.com/layerfMRI/Sequence_Github/ tree/master/3T.

3.2 Cerebrovascular reactivity

Cerebrovascular reactivity (CVR) reflects the capacity of brain vessels to dilate or constrict in response to a vasoactive stimulus. MRI-based CVR is generally expressed as percent BOLD or ASL signal change per mmHg change in end-tidal carbon dioxide ($EtCO_2$) [2]. Despite the relative simplicity of the measure, it is one of the most reliable MRI-derived predictors of cerebrovascular impairment and has been applied in the study of several diseases, including SVD and AD [2, 35, 36].

Since GRE-BOLD signal changes are heavily weighted towards the venous macrovasculature, both due to an inherent weighting towards larger vessels and to dHb-containing compartments [37], CVR maps based on such contrast show similar features. In order to target microvascular CVR, different sequences should be employed, such as ASL, spin-echo (SE)-BOLD, or VASO. In particular, VASObased CVR could be interesting due to the high resolutions achievable and the observed stability to hypercapnic stimuli [14]. In one study, VASO reactivity was found to be more negative (i.e., larger CBV increase) in patients with carotid artery disease compared to healthy controls [38].

3.2.1 Vasodilatory stimuli

CVR mapping with MRI can be achieved with administration of carbon dioxide (CO₂)-enriched gas [2], breath holding [39] or amplitude of resting-state fluctuations in CO₂ [40]. CVR mapping has been performed at a variety of spatial resolutions, determined by the BOLD or ASL acquisition protocol used. For CVR mapping at high resolution, inhalation of a CO₂-enriched gas mixture (typically 5% CO₂) is preferable over the other approaches because it gives the strongest increase in CBF and thus improves the contrast-to-noise ratio (CNR), while being safe

and well tolerated [41]. A few studies have indeed successfully mapped CVR at voxel sizes below 2 mm using breathing manipulations [2, 42]. In particular, a recent study by Schellekens et al. reports depth-dependent profiles of CVR using GRE-BOLD and SE-BOLD, showing a detectable variation with cortical depth [15].

3.3 Cerebral metabolic rate of oxygen consumption

BOLD signal changes in a voxel during a task of interest can be modeled according to Eq. 1 [4]:

$$\frac{\Delta \text{BOLD}}{\text{BOLD}_0} = M \left[1 - \left(\frac{\text{CBV}_{\nu}}{\text{CBV}_{\nu,0}} \right) \left(\frac{\text{CBF}}{\text{CBF}_0} \right)^{-\beta} \left(\frac{\text{CMRO}_2}{\text{CMRO}_{2,0}} \right)^{\beta} \right]$$
(1)

where CBV_v is the CBV contributing to BOLD signal changes ("venous" CBV), *M* is the calibration parameter which corresponds to the maximum BOLD signal change (BOLD = *M* for complete deoxy-hemoglobin washout from the vasculature), β is a magnetic-field dependent exponent, and the subscript 0 indicates quantities at baseline.

It is generally assumed that CBV and CBF are coupled via Eq. 2:

$$\frac{\text{CBV}_{\nu}}{\text{CBV}_{\nu,0}} = \left(\frac{\text{CBF}}{\text{CBF}_0}\right)^{\alpha} \tag{2}$$

where α is a coupling constant, often taken to be equal to 0.38 [43].

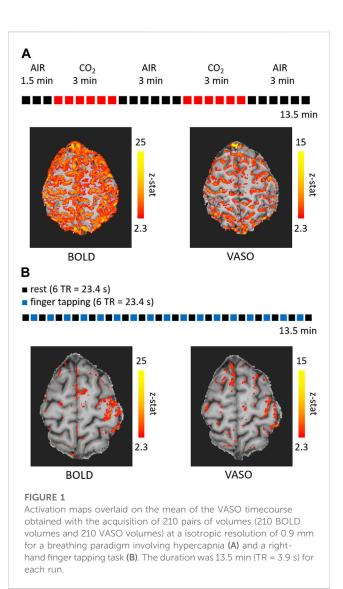
The goal of calibrated fMRI is isolating the CMRO₂ term from the Eq. 1: to this end, a pure CBF or CBV sensitive contrast needs to be acquired in addition to BOLD, and the calibration parameter *M* needs to be estimated. For extracting *M* a vasoactive, isometabolic (CMRO₂/CMRO_{2,0} ~ 1) stimulus is used, which introduces a BOLD dependency only on CBF or CBV (measurable) if Eq. 2 is assumed [4, 5].

Most calibrated fMRI studies use ASL sequences for mapping CBF and extract CBV via Eq. 2. Since the technique allows for voxelwise mapping, the resolution achievable by ASL at 3 T dictates the resolution of CMRO₂ maps. For submillimetric applications, CBV mapping can be used and CBF changes estimated by Eq. 2, while assuming a different coupling exponent for venous and total CBV changes [14, 15].

Other approaches for $CMRO_2$ mapping have been introduced using different gas mixtures or without exogenous gas manipulations. Existing acquisition techniques and modelling approaches have been recently reviewed [44, 45].

4 Functional paradigms for multiparametric mapping at high resolution

Given the sequences currently available, we propose that it is possible to obtain, in less than 30 min at 3 T, submillimetric maps of 1) BOLD-based CVR; 2) VASO-based CVR; 3) the calibration parameter M; 4) relative changes in CMRO₂ elicited by a task of interest; 5) BOLD signal changes elicited by the same task; 6) VASO signal changes elicited by the same task.



To this end, the use of the VASO sequence with a 3D readout [13] has shown to be suitable for submillimetric fMRI at 3 T.

Figure 1 shows the activation maps obtained for two tasks using the same acquisition protocol. The sequence used was the 3 T VASO [13] with the following acquisition parameters: isotropic nominal resolution = 0.9 mm, matrix size = $188 \times$ 188×28 , echo time (TE) = 21.2 ms, TR = 3.9 s (this refers to the time it takes to acquire a BOLD-VASO pair of volumes), 210 measurements (210 BOLD volumes and 210 VASO volumes), partial Fourier = 6/8, GRAPPA = 3, bandwidth = 1,026 Hz/Px, echo spacing = 1.1 ms, inversion delay = 550 ms, flip angle = 30° with a variable flip angle scheme [13], water selective excitation achieved with long binomial 1-1 pulses. The acquisition was performed on a Siemens Prisma 3 T (Siemens Healthineers, Erlangen, Germany) scanner and a 32-channel head coil. Informed consent was obtained according to international standards, the study was approved by the Ethical Committee of Fondazione Santa Lucia, CE/2022_010.

One acquisition was performed during a breathing challenge consisting of an alternation of medical air and 5%-CO₂-enriched air administration (Figure 1A), for a total acquisition time of

13.5 min. The other acquisition was performed during a functional task consisting of an alternation of right-hand finger tapping for 6 TRs and rest for 6 TRs (Figure 1B). For both acquisitions, the timecourses were corrected for motion using the ASL toolbox based on SPM [46, 47]. Statistical activation maps were obtained using FSL FEAT [48] and z-stats are reported in Figure 1. A detectable activation was present for both contrasts and conditions, with BOLD generating larger clusters as expected. The submillimetric voxel size of 0.9 mm allows to have 2 to 4 voxels within the gray matter thickness and to reduce partial voluming with the neighboring CSF and WM. Nevertheless, the large BOLD signal changes following the pial venous vasculature are located outside the GM and reduce the spatial specificity. VASO signal changes have in general a lower amplitude and are more affected by noise, which result in smaller clusters as in Figure 1, while the specificity of the signal changes is less compromised by the pial vasculature (more clearly visible in Figure 1B).

This preliminary dataset confirms that it is possible to have a statistically significant response in both BOLD and VASO timecourses for a functional run of 13.5 min (210 BOLD-VASO pairs of volumes).

With the two datasets referred to in Figure 1, the signal changes in BOLD and VASO upon hypercapnia can be used to estimate the calibration parameter M across gray matter from Eq. 1 (assuming Eq. 2), and the signal changes in BOLD and VASO elicited by the finger tapping task can then be used to estimate changes in CMRO₂ from the same equation, once M has been calculated.

4.1 Limitations

The proposed multiparametric acquisition can offer a multifaceted window into microvascular dysfunction, by mapping vascular and metabolic features concomitantly at 3 T and at voxel volumes below 1 mm³.

The submillimetric VASO sequence has been introduced only recently at 3 T and has some limitations. First of all, the low SNR of 10-20 [13] limits its flexibility: tasks associated with small effect sizes are hard to detect, unless the acquisition time is largely increased. Moreover, the generation of the VASO contrast has some timing constraints that impact the TR duration: the typical TR is about 4-5 s (for the acquisition of a full BOLD-VASO pair) for slab protocols.

Additionally, the breathing challenge could create excessive discomfort in patient populations, both due to wearing a mask and being in a hypercapnic state. Therefore, it is important to familiarize each subject with the gas challenge prior to scanning. An earlier study showed an increase in participant dropouts for AD patients compared to healthy control for the breathing challenge [26].

While CVR mapping is reliable and has been applied in disease, the same does not hold for calibrated fMRI. Calibrated BOLD models rely on several assumptions, which, even at typical image resolutions, might not be a good approximation of the real underlying mechanisms. Such limitations could be exacerbated at higher resolutions where the spatial heterogeneity of the vasculature is resolved. Most notably, the flow-volume coupling expressed by Grubb's relationship is unlikely to hold true with the same exponent for small voxel sizes and between healthy subjects and patients. Therefore, differences in CMRO₂ between a healthy group and a disease group could stem from a disrupted CBF-CBV coupling rather than a true metabolic difference. To avoid such bias, imaging CBF, CBV, and BOLD concomitantly [8] is preferable, but CBF mapping via arterial spin labeling at high resolution is challenging [49], therefore studying the flow-volume coupling relationship experimentally in humans is difficult.

If using VASO in the calibrated BOLD framework, the CBV at baseline needs to be assumed. A fixed fraction of 5.5% [50] might be acceptable considering that the microvascular blood density, which is the major factor responsible for VASO signal changes for short stimulations, is relatively homogeneous across the cortical depth [10, 51]. During hypercapnic stimulations, however, the fraction of VASO signal change coming from venous blood increases [52], challenging the assumption of a fixed baseline blood fraction.

Finally, the assumption of isometabolism during gas administration has been challenged in several studies. Although there is not a definitive answer, some evidence suggests a reduction, rather than an increase, in CMRO₂ of up to 13% during a 5%-CO₂ gas challenge [53]. However, this value depends on the concentration of arterial CO₂ and on the length of the respiratory challenge, with shorter epochs having less impact on CMRO₂ [2].

5 Discussion and conclusion

We propose that, to obtain more insight into neurovascular diseases, several contrasts and approaches sensitive to microvascular function should be used at 3 T and at submillimetric resolutions. The acquisition scheme that we propose here involves the use of VASO at 3 T (BOLD and CBV-weighted contrast interleaved) with a gas manipulation inducing hypercapnia, and a functional task. Scanning under these conditions for less than 30 min makes it feasible to obtain maps of BOLD and VASO signal changes for each condition, CVR, regional CMRO₂ changes and M at a submillimetric resolution.

Data availability statement

The raw data supporting the conclusion of this article will be made available by the authors, without undue reservation.

Ethics statement

The studies involving humans were approved by the Ethical Committee of Fondazione Santa Lucia. The studies were conducted in accordance with the local legislation and institutional requirements. The participants provided their written informed consent to participate in this study. Written informed consent was obtained from the individual(s) for the publication of any potentially identifiable images or data included in this article.

Author contributions

MG: Conceptualization, data collection and analysis, writing. GG: Conceptualization, data analysis, writing. EB, RW, and FG: Conceptualization, writing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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