



Clinician Accessible Tools for GUI Computational Models of Transcranial Electrical Stimulation: BONSAI and SPHERES



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ABSTRACT

Computational models of brain current flow during transcranial electrical stimulation (tES), including transcranial direct current stimulation (tDCS) and transcranial alternating current stimulation (tACS), are increasingly used to understand and optimize clinical trials. We propose that broad dissemination requires a simple graphical user interface (GUI) software that allows users to explore and design montages in real-time, based on their own clinical/experimental experience and objectives. We introduce two complimentary open-source platforms for this purpose: BONSAI and SPHERES. BONSAI is a web (cloud) based application (available at neuralengr.com/bonsai) that can be accessed through any flash-supported browser interface. SPHERES (available at neuralengr.com/spheres) is a stand-alone GUI application that allow consideration of arbitrary montages on a concentric sphere model by leveraging an analytical solution. These open-source tES modeling platforms are designed go be upgraded and enhanced. Trade-offs between open-access approaches that balance ease of access, speed, and flexibility are discussed.

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Broad need but limited access to tES (tDCS) computational models

Computational “forward” models predict brain current flow during tDCS [1,2] as well as during other non-invasive transcranial electrical stimulation (tES) techniques [3] such as CES [4], tACS [5]. Because the relationship between stimulation dose (defined as those electrode and waveform parameters controlled by the operator [6]) and resulting brain flow is complex and non-intuitive [7], computational forward models underpin how protocols are designed and understood. Though model validation efforts are ongoing [8,9], these models represent a standard to understand brain current flow and optimize tES dose and so inform clinical practice and behavior research on an ongoing basis [10–12].

Yet despite increased interest in tES modeling, as supported by the number of tES publications about or including a modeling component (from 82 per year in 2009 when we proposed the first

gyri-precise model to >359 per year expected for 2013), access to modeling tools by clinicians remains highly limited. And ironically much of the effort to enhance the relevance of modeling through increases sophistication (complexity [13]) in fact hinders both reproduction and dissemination. Efforts to automate a modeling “pipeline” still require some engineering/computational proficiency [14,15]. Though several recent effort have compartmentalized the tools needed [16,17].

We propose that broad dissemination requires an intuitive, simple, and stand-alone software. Moreover, the ability to user to explore and design montages in real-time, based on their own clinical/experimental experience and objectives, is critical to develop intuition on dose strategies [18]. This is because decision about what dose and related brain current flow pattern is “optimal” will be dependent on the specific study objective and investigator bias on neuromodulation mechanisms (e.g. role of collateral targets). Here we introduce two complimentary open-source platforms for this purpose: SPHERES and BONSAI.

Fundamental challenge in dissemination

Any attempt to enhance access for clinicians and behavior researchers to tES modeling must address two fundamental

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challenges. First and foremost is the myriad of potential montages that can be evaluated (variations in electrode number and for each electrode: waveform, current, size, position, and shape [6]) compounded by individual variations in current flow for any given montage [3,19]. Inevitably, modeling publications on dose design can consider only a very limited selection of montages, and one or a limited set of “representative” head models. Pipeline computational processes accelerate this process but do not remove the underlying problem. Ironically, it is precisely because models illustrate the importance of montage and anatomy, that generic models may be imprecise. Our recent invention leveraging linearity allows for an approach where individual High-definition electrode activation are pre-solved for a given head model, and then any HD and conventional pad montage (by grouping adjacent electrodes) is instantly predicted by linear summation [20]; moreover an optimal montage can be predicated for any given target and orientation [21] without the need for iterative search. None-the-less, these new approaches do not remove the need for extensive computational resource for pre-processing, are individual head specific, and are further limited by assumptions on tissue segmentation and tissue conductivity [22].

The second fundamental challenge relates to representation of brain current flow. Most modeling studies follow the “quasi-uniform” assumption and presume neuromodulation is represented by local tissue current density or electric field [23]. Even so, the

combination of multiple regions being stimulated (a large fraction of the entire brain for conventional tDCS montages [24]) compounded by the details of idiosyncratic current flow (changing even with a single gyri [25]) make representation complicated. Even for a given montage and head, the regions and details of interest will vary across investigators. Inevitably, publications cannot with limited figures detail brain current flow in every region of potential interest.

The SPHERES and BONSAI software each address these limitations in a distinct manner while meeting our requirement that modeling tools be instantly and simply accessible. The overall objective is to encourage exploration, and analysis, and optimization of tES dose by the behavioral scientists and clinicians.

SPHERES

SPHERES (available at neuralengr.com/spheres) is a stand-alone graphical user interface (GUI) application that allow consideration of arbitrary montages on a concentric sphere model by leveraging an analytical solution [26]. SPHERES further allows adjustment of “tissue” parameters, namely sphere thickness and conductivity. The technique is rooted in the spherical harmonic expansion of the applied scalp currents and induced electric fields, allowing for a linear systems formulation of the TES forward problem. SPHERES

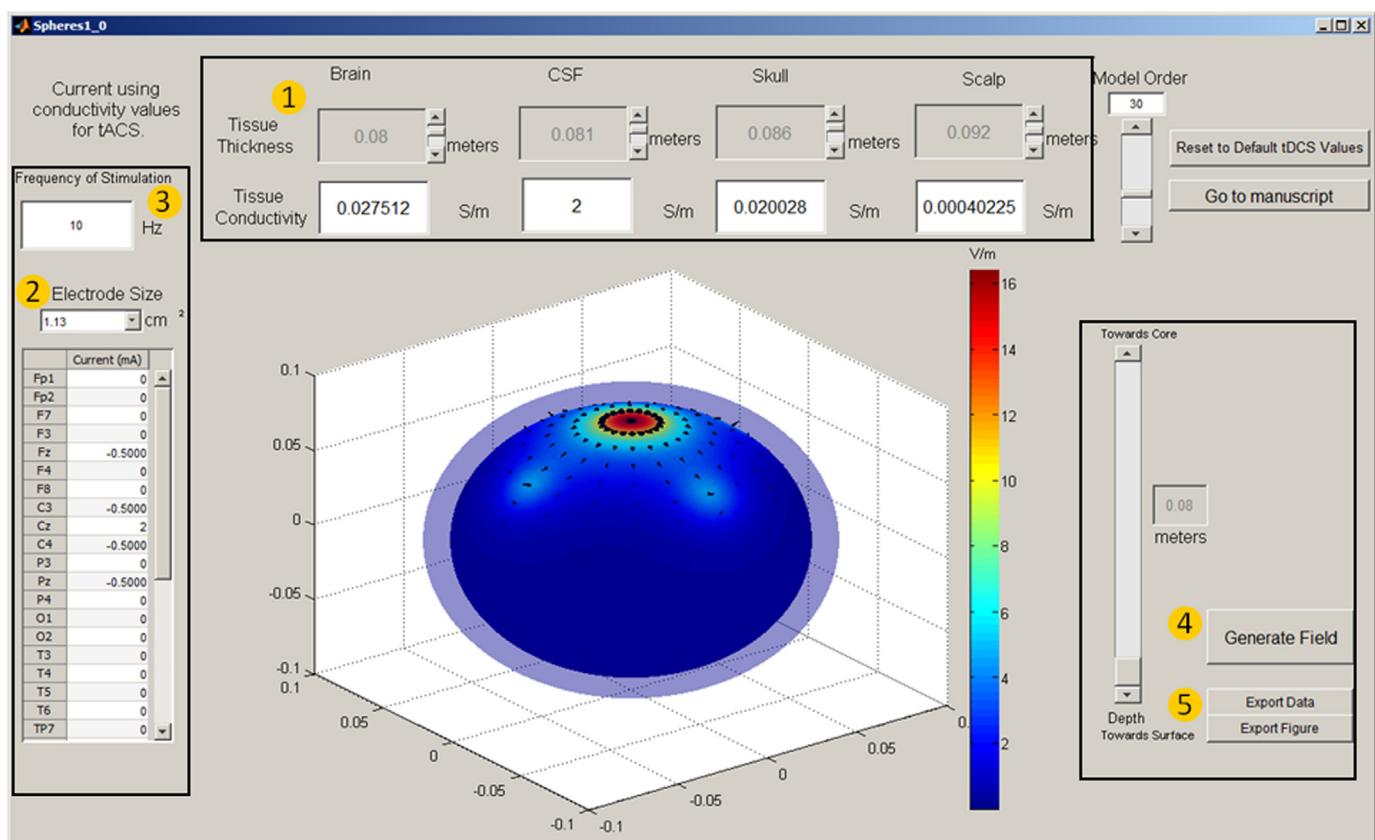


Figure 1. SPHERES modeling software allows simple and rapid simulation in a concentric spheres head representation (available free at neuralengr.com/spheres) (1). Head parameters: Users are able to adjust the tissue thickness for brain, CSF, skull, and scalp with their respective slider in meters, as well as entering the corresponding conductivity values for each tissue (2). Electrode parameters: Users are able to select the size of the electrode being used in the drop down menu, and enter the amount of current being applied in mA at the specific positions. Note the anode current positive while the cathode current is negative, the sum of the currents entered in the table should add up to zero (3). Entering a frequency of stimulation value will automatically change the tissue conductivities to tACS values. To return to tDCS values, the user needs to press “Reset to Default Values” as tDCS is the default option (4). “Generate Field” button will compute your input values and generate the corresponding figure. The movement of the slider changes how deep you are looking inside the human head. The current depth is displayed in meters (5). “Export Data” will allow the users to save the data used to generate the current image in a “.mat” file, only usable by matlab. “Export Figure” will directly allow users to save the figure.

can reproduce and expand on any numerical simulation study on tES (Fig. 1).

At a time when increasing complexity often drives modeling efforts, the rationale for concentric sphere simulations should be emphasized. It is *precisely* because head anatomy is irregular, and indeed highly individual, that sphere modeling provides an initial basis to consider the principles of dose design [1,13,27,28]. For example, using spheres the role of inter-electrode distance or CSF conductivity can be considered independent, while in a realistic head multiple confounds from other anatomical factors (e.g. inhomogeneous skull thickness under the electrode, proximity for foramen [29,30]) cannot be excluded. Spherical models ignore the critical role of tissue inhomogeneity and anatomical detail, especially cortical folding [24,25,31]. None-the-less, spherical models are useful to understand what features of dose design and tissue properties are thus valuable tool to both experts and novices.

In regards to the first challenge to dissemination, SPHERES uses a closed-form analytical solution to allow modeling of any arbitrary electrode montage (the “point spread function of the

head” [26]) with little computation (seconds on a conventional PC); moreover, tissue properties, including frequency-specific tissue resistivity, can be readily adjusted. Inter-individual variation can be considered in the abstract sense by individual sphere radius or conductivity [32]. In regards to the second challenge of dissemination (for better or for worse), it is straightforward to represent current flow through the spheres precisely because anatomical detail is absent – though quantitative inferences about current spread (focality and deep structures) should be made with caution. The overall result of the SPHERES program is a tool-box that allows infinite variations in dose and analysis by users, specifically for their given application, consistent with the overall objective.

BONSAI

BONSAI is a web (cloud) based application (available at neuralengr.com/bonsai) that can be accessed through any flash-supported browser interface. Data from any simulation of current

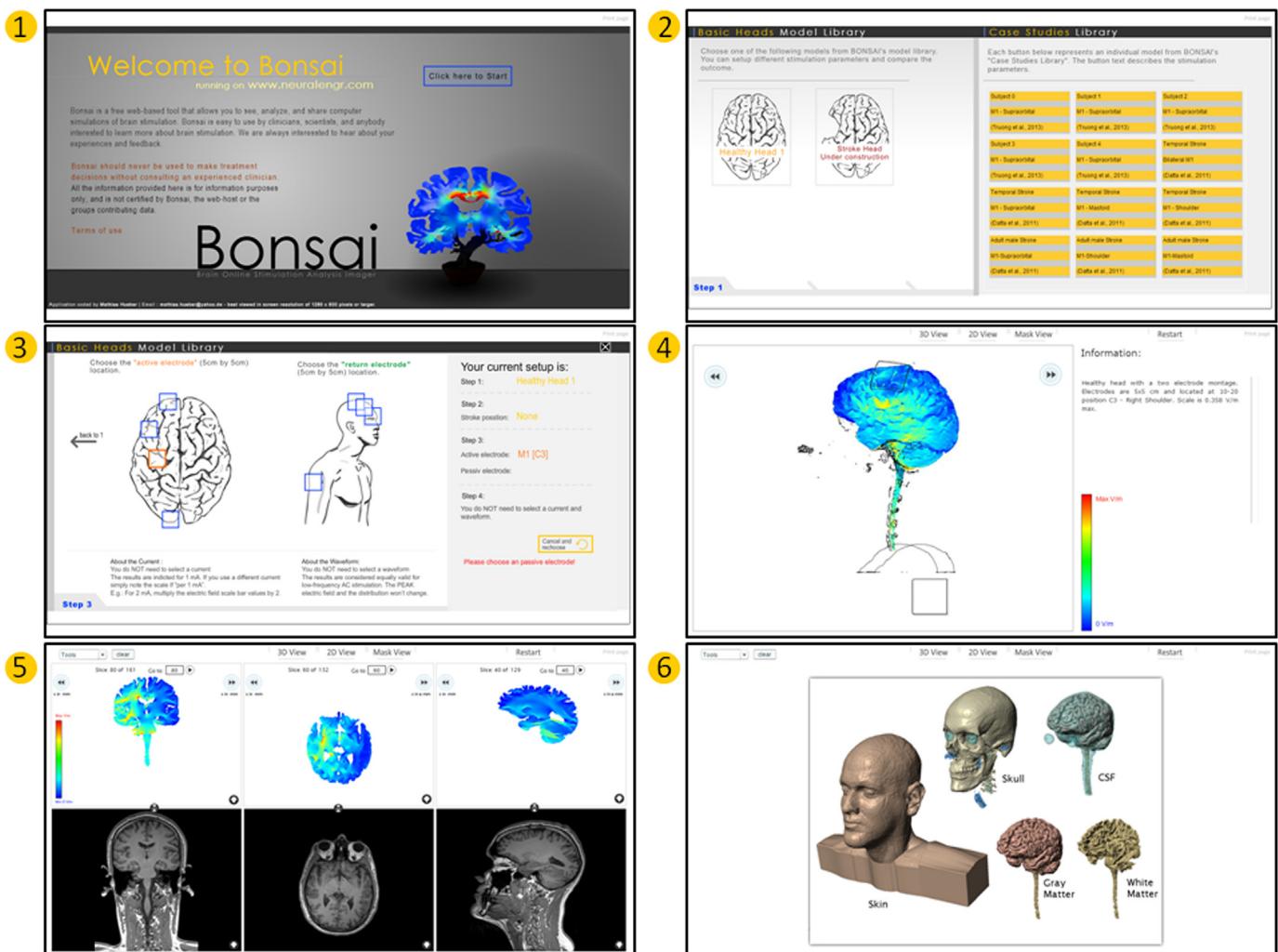


Figure 2. BONSAI modeling software allows selection from pre-solved high-resolution MRI-derived models simulations and web-based exploration of current flow (available free at neuralengr.com/bonsai). Bonsai user interface walkthrough: Click start (1). Select a “Basic Head” or a “Case Study” (2). If a “Basic Head” was selected, now select the electrode configuration. Left click a box under “active electrode” and under “return electrode,” and then click “Load the data now” (3). After loading, a 3-D rendering of the electric field intensity can be rotated left or right by clicking the respective arrows or by clicking and dragging over the image. Information about the electrode configuration and intensity scale are given on the right (4). Alternate views of the data, 2-D slices (5) or a rendering of the segmentation (6), can be selected from the tabs at the top of the window. Under “2-D View” coronal, axial, and sagittal slices of the electric field intensity are displayed along with their respective MRI slice. The images can be overlaid by clicking the converging arrows between the FEM solution and MRI. Pen/highlighting tools are available at the top left (5).

flow, regardless of how it was generated, can be uploaded as serial images and then viewed using the BONSAI web interface. Evidently, users cannot consider changes in montage of head models not already included (uploaded) — BONSAI does not support, for example, adjusting the position of an electrode to an arbitrary location. To automatically parse data from commercial FEM solvers to BONSAI scripts may be developed. Regardless of FEM package, solutions can be exported in a text format as a list of nodes and elements with corresponding electric field values. Scripts in software packages like Matlab or Python can be used to interpolate, scale, and export slice images co-registered to the original model MRI. If the data were derived from a given publication, that publication is referenced both providing insight into the given approach and enhancing the impact of the publication. If a suitable approximation of simulation montage is available in BONSAI, it can be used to inform dose design and in subsequent publications (Fig. 2).

In regards to the first and second challenge of dissemination BONSAI takes a highly accessible and simple (if “brute force”) approach of simply coalescing individual modeling efforts in indexed images. However, in contrast to dispersed individual publications, 1) BONSAI represents a growing database on montages and approaches that can be compared and contrasted, 2) the methods and results are fully disclosed allowing reproduction and further analysis; 3) and the BONSAI interface allows sharing of more images (cross-sections) than possible in any publication. Though limited in flexibility, the overall result of the BONSAI “library” is a tool-box that encourages analysis and optimization by users for their given application consistent with the overall objective of dissemination.

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