

1 Effect of field spread on resting-state MEG functional network
2 analysis: a computational modeling study

3 Silva Pereira¹, S., Hindriks¹, R., Mühlberg¹, S., Maris², E., van Ede², F.,
4 Griffa^{3,4}, A., Hagmann^{3,4}, P., Deco^{1,5}, G.

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5 1. Center for Brain and Cognition, Computational Neuroscience Group, Department of Information
6 and Communication Technologies, Universitat Pompeu Fabra, Barcelona, Spain.

7
8 2. Radboud University Nijmegen, Nijmegen, The Netherlands.

9
10 3. Department of Radiology, Lausanne University Hospital and University of Lausanne, 1011 Lau-
11 sanne, Switzerland.

12
13 4. Signal Processing Laboratory 5, Ecole Polytechnique Federale de Lausanne, 1015 Lausanne, Switzer-
14 land.

15
16 5. Institució Catalana de la Recerca i Estudis Avanats (ICREA), Universitat Pompeu Fabra, Barcelona,
17 Spain.

Abstract

19 A popular way to analyze resting-state EEG and MEG data is to treat them as a functional network
20 in which sensors are identified with nodes and the interaction between channel time-series with the
21 network connections. Although conceptually appealing, the network-theoretical approach to sensor-
22 level EEG and MEG data is challenged by the fact that EEG and MEG time-series are mixtures of
23 source activity. It is therefore of interest to assess the relationship between functional networks of
24 source activity and the ensuing sensor-level networks. Since these topological features are of high
25 interest in experimental studies, we address the question of to what extent the network topology can
26 be reconstructed from sensor-level FC measures in case of MEG data. Simple simulations that consider
27 only a small number of regions do not allow to assess network properties; therefore, we use a diffusion
28 MRI-constrained whole-brain computational model of resting-state activity. Our motivation lies behind
29 the fact that still many contributions found in the literature perform network analysis at sensor level,
30 and we aim at showing the discrepancies between source- and sensor-level network topologies using
31 realistic simulations of resting-state cortical activity. Our main findings are that the effect of field
32 spread on network topology depends on the type of interaction (instantaneous or lagged) and leads
33 to an underestimation of lagged FC at sensor level due to instantaneous mixing of cortical signals,
34 instantaneous interaction is more sensitive to field spread than lagged interaction, and discrepancies
35 are reduced when using planar gradiometers rather than axial gradiometers. We therefore recommend
36 to use lagged interaction measures on planar gradiometer data when investigating network properties
37 of resting-state sensor-level MEG data.

1 Introduction

Electroencephalography (EEG) and magnetoencephalography (MEG) allow for non-invasive monitoring of electrical activity in the human brain at a time-scale of milliseconds [21]. EEG and MEG have been used extensively to characterize functional connectivity (FC) underlying healthy and impaired cognitive processes, and promoted the development of methods for characterizing FC in electrophysiological recordings (EEG/MEG). Today, one can choose from a wide variety of FC measures based on different mathematical theories, such as chaotic systems, weakly-coupled oscillators, linear stochastic processes, and information theory [36, 40, 44]. In applications to EEG and MEG data, it is common practice to calculate the FC measure for all channel-pairs, resulting in an FC matrix with the corresponding one-to-one relationships. The obtained FC matrices of different subjects are subsequently analyzed and can be used to characterize different cognitive states or pathological conditions. In this study, we focus on the fundamental question of what it is that sensor-level FC matrices say about the dynamical structure of the data, and to what extent the network topology can be reconstructed from sensor-level FC measures in case of MEG data. To that end, we compare sensor-level FC matrices with the underlying cortical FC matrices.

The most common way to think about EEG/MEG FC matrices is in terms of functional networks [8, 23, 39]. Viewed as such, network nodes are assumed to correspond to cortical circuits underlying the EEG/MEG channels and network connections are assumed to correspond to functional dependencies between these circuits. Analysis of the network is typically carried out using graph-theoretic measures such as the clustering coefficient, degree distribution, and average path length. The network approach to EEG/MEG brain dynamics has demonstrated that resting-state functional networks at sensor level possess small-world properties in several frequency bands [1, 46], which would enable fast and robust integration and segregation of information [2]. Moreover, graph-theoretic measures are able to discriminate between the ongoing brain dynamics of healthy individuals and those of psychiatric and neurological patients [29, 30, 37, 38].

Although these results are encouraging and clinically relevant, by themselves they do not validate the network-theoretic view on sensor-level EEG/MEG brain dynamics. In addition, the network approach to EEG/MEG brain dynamics is questionable from a physical perspective. Concerning the physics, because EEG/MEG channels register source activity not only from tissue directly underlying the channels but from distant sources as well, they cannot directly be associated with nodes of a functional network and the FC measures suffer from spurious correlations due to the superposition of sources [22]. Moreover, scalp-based EEG connectivity estimates like vector autoregressive models have been shown to be highly affected by volume conduction [7]. To isolate true FC from spurious FC due to volume-conduction (EEG) or field spread (MEG), several approaches have been proposed. Since volume-conduction/field spread is instantaneous, an alternative is to focus on time-lagged interactions between EEG/MEG channels [26, 32, 33, 47], which necessarily reflect time-lagged interactions between sources. However, although these FC measures reduce spurious correlations and might lead to more faithful networks [5, 19, 45], it remains unclear whether such measures actually reflect functional interaction between local circuits. Another alternative is to perform a source space projection prior to calculation. Source reconstruction techniques like beamforming have become standard for source space projection of EEG/MEG sensor data, and recent studies have proposed methods to mitigate the effects of volume-conduction/field spread [10, 18, 41, 49]. Although these techniques have been shown to ease the effects of field spread, still several contributions continue applying network theory to sensor level data, and in this study we highlight the consequences of carrying out such an analysis.

Motivated by the existence of several fundamental methodological difficulties [22], we use models based on empirical data to uncover a discrepancy between cortical FC matrices and those computed at channel level. Whereas the authors in [22] focused on a single minimalistic scenario, the authors

89 in [10] studied the performance of a novel connectivity measure on both simulated and real data at
90 source-level using simple networks composed of only a few nodes (five), without performing realistic
91 simulations of resting-state cortical activity and without reporting results on sensor-level connectivity
92 studies. The extended analysis we perform on simulated data can be contrasted with those based on a
93 network view on EEG/MEG brain dynamics, e.g. [6, 15, 16, 29, 45, 46]. To this extent, we focus on both
94 real and complex-valued correlation measurements (or coherence), and argue that treating sensor-level
95 resting-state EEG/MEG recordings as comprising a discrete network of interacting systems might not
96 be appropriate and obscures rather than clarifies the true brain dynamics, since after all, the recorded
97 electric and magnetic fields are continuous in space [42]. To demonstrate this, we use a diffusion MRI-
98 constrained whole-brain computational model of resting-state activity, where the activity at different
99 areas of the brain emerges through long-range synchronization of Hopf oscillators. We stress that in
100 contrast to existing simulation studies on sensor-level FC which typically use simple network models,
101 we perform realistic simulations based on empirical data. The measured MEG signals are obtained
102 from a forward model assuming two different systems: sensors with axial gradiometers and sensors
103 with planar gradiometers, the latter obtained synthetically using the Fieldtrip toolbox [35]. Then,
104 we perform a network analysis at both source level and sensor level, and contrast the results showing
105 that they can lead to wrong interpretations. A Louvain modularity study is included, which sheds
106 additional light on the results from the standard network analyses performed.

107 **2 Materials and Methods**

108 **2.1 Cortical parcellation and white-matter tractography**

109 For the estimation of the whole-brain structural connectivity matrix S , we considered ten male healthy
110 subjects aged 22.5 ± 1.8 years, nine right-handed and one left-handed. Each subject underwent a
111 magnetic resonance imaging (MRI) session on a 3 Tesla scanner (Magnetom TrioTim, Siemens Medical
112 Solutions), equipped with a 32-channel head coil. Each MRI session included a magnetization-prepared
113 rapid acquisition gradient echo (MPRAGE) and a diffusion spectrum imaging (DSI [52]) sequence.
114 The MPRAGE acquisition had 1 mm in-plane resolution and 1.2 mm slice thickness, covering $240 \times$
115 257×160 voxels. The TR, TE and TI were 2300, 2.98 and 900 ms. The DSI acquisition had $2.2 \times$
116 2.2×3 mm resolution, covering $96 \times 96 \times 34$ voxels, and it included 256 diffusion-weighted volumes
117 (with maximum b-value 8000 s/mm^2) and one b0 volume. The TR and TE were 6100 and 144 ms.
118 Informed written consent in accordance with institutional guidelines was obtained for all participants.
119 Subject-wise MPRAGE volumes were segmented into white-matter, grey-matter and cerebrospinal-
120 fluid compartments using FreeSurfer software [12], and linearly registered to the diffusion space (b0
121 volume) [20]. The grey-matter cortical volumes were segmented into 219 regions of interest (ROIs),
122 according to the atlas described in [9] and using the Connectome Mapper Toolkit [11]. DSI data
123 were reconstructed according to [52]. Deterministic streamline tractography [27] was performed on
124 reconstructed DSI data, initiating 32 streamline propagations per voxel and per diffusion direction. A
125 representative structural connectivity matrix S was estimated by combining tractography and cortical
126 parcellation results from the ten subjects.

127 **2.2 Computational model of resting-state cortical dynamics**

128 Recently, it has been reported that the dynamics of resting-state cortical fluctuations is poised at
129 the transition between asynchronous to oscillatory behavior [14]. We use the model proposed in [14]
130 to simulate resting-state cortical activity, where a general neural mass model based on the normal
131 form of a Hopf bifurcation was proposed. The model describes the dynamics of n coupled brain areas
132 (nodes). The global dynamics emerges through the mutual interactions between the nodes, which
133 are coupled through a connectivity matrix S . The local dynamics are modeled by the normal form
134 of a supercritical Hopf bifurcation, which describes the transition from asynchronous noisy behavior

135 to self-sustained oscillations, as controlled by the bifurcation parameter, denoted a . The complex-
 136 valued cortical activity at every node j is modeled by the following set of coupled non-linear stochastic
 137 differential equations:

$$\frac{d}{dt}z_j(t) = (a + i\omega_0)z_j(t) - z_j(t)|z_j(t)|^2 + g \sum_{k=1}^n S_{j,k}(z_k(t - t_{j,k}) - z_j(t)) + \eta_j(t), \quad (1)$$

138 where i is the imaginary unit, ω_0 is the intrinsic frequency, η_j is additive Gaussian white noise with
 139 intensity σ_j , and $t_{j,k}$ denotes the axonal delay between node j and node k . The parameter g globally
 140 scales the synaptic connections and is used to control the correlation between regions. When the nodes
 141 are uncoupled (i.e., $g = 0$), the local dynamics bifurcate when a crosses zero. Specifically, for $a < 0$, the
 142 local dynamics have a stable fixed point at $z_j = 0$, which corresponds to a low-activity asynchronous
 143 state due to the additive noise. For $a > 0$, there exists a stable limit-cycle with angular frequency ω_0 .
 144 The effect of the bifurcation parameter value a on the local dynamics is illustrated in Fig. 1.

145
 146 The complex-valued variable z_j can be decomposed into a real part x_j and imaginary part y_j as
 147 $z_j = x_j + iy_j$. In terms of x_j and y_j , Eq. (1) takes the form

$$\frac{d}{dt}x_j = ax_j - \omega_0 y_j - x_j(x_j^2 + y_j^2) + g \sum_{k=1}^n S_c(j,k)(x_k - x_j) + \eta_j, \quad (2)$$

$$\frac{d}{dt}y_j = ay_j - \omega_0 x_j - y_j(x_j^2 + y_j^2) + g \sum_{k=1}^n S_c(j,k)(y_k - y_j) + \eta_j, \quad (3)$$

148 where we have suppressed the time-dependence of x and y . Cortical activity at node j is thus modeled
 149 by the real-valued variable x_j .

150
 151 To simulate realistic cortical resting-state activity, the nodes of the computational model are taken to
 152 be the ROIs in the surface-based cortical atlas described in Sec. 2.1, and the connectivity matrix is
 153 taken to be the group-averaged structural connectivity matrix obtained from diffusion MRI tractog-
 154 raphy. To obtained activity in each vertex of the cortical mesh, the activity for each ROI is copied to
 155 all vertices within that ROI. This effectively means that we assume cortical activity to be locally syn-
 156 chronous over several squared centimeters, which is consistent with empirical estimates of the spatial
 157 extent of cortical activity underlying resting-state EEG and MEG recordings [21, 24, 34].

158
 159 From the simulated cortical activity, MEG sensor data was computed by constructing a leadfield
 160 matrix for the 275-channel MEG gradiometer system (CTF Systems Inc., Port Coquitlam, Canada)
 161 using a single-sphere head model in the Fieldtrip toolbox [35]. The source space was restricted to the
 162 vertices of the left and right Freesurfer cortical template meshes, which were manually positioned inside
 163 the MEG helmet. The single-sphere had a radius of 10 cm and was manually positioned to enclose the
 164 source space. Dipoles were assumed to be oriented perpendicular to the local cortical surface. This
 165 yielded leadfield matrices of dimensions 273×130385 for the left hemisphere and 273×131547 for the
 166 right hemisphere (the system has two reference channels not used for recording). Besides these axial
 167 gradiometer data, we computed the corresponding planar gradiometer data using Fieldtrip [35].

168
 169 Noisy MEG data is obtained by adding correlated Gaussian noise to the simulated signals, for dif-
 170 ferent values of the peak signal-to-noise ratio (PSNR), given by

$$PSNR = 20 \log \frac{\max_{i,t} |s_i(t)|}{\sigma_N}, \quad (4)$$

171 where $s_i(t)$ is the noise-free MEG signal at channel i at time t . The noise covariance matrix was
 172 obtained from an empty-room MEG recording and further scaled by a factor $\sigma_N^2/\bar{\sigma}^2$ to obtain the
 173 desired PSNR values, where $\bar{\sigma}^2$ is the mean of the entries of its main diagonal.

2.3 Functional connectivity matrices

There exist a wide range of measures to characterize functional connectivity between two simultaneously recorded time-series [36] and most of them have been applied to sensor-level EEG and MEG recordings [40]. Since EEG and MEG sensor-level time-series are linear superpositions of cortical time-series (volume-conduction/field spread), the results of sensor-level FC analyses are generally difficult to interpret. This motivated the development of FC measures that are based on time-lagged interactions, since these cannot be explained by field spread. Examples of such measures are imaginary coherency [33], phase-slope-index [32], phase-lag-index [47], and the (symmetric part of the) phase-modulation function [26]. In the current study we consider two such measures: the imaginary part of the complex-valued Pearson correlation coefficient (IC) and the phase-lag index (PLI). For comparison, we also use the real part of the (complex-valued) Pearson correlation, which quantifies instantaneous FC. Below, a short description of these three measures is given.

Given two oscillatory time-series $x = (x_1, \dots, x_n)$ and $y = (y_1, \dots, y_n)$, the *complex-valued Pearson correlation coefficient* $r_{x,y}$ between x and y is defined as the Pearson correlation coefficient between the analytic signals x^A and y^A of x and y , respectively. Thus, let $x^A = x + ix^H$ and $y^A = y + iy^H$ denote the analytic signals of x and y , respectively, where x^H and y^H denote the Hilbert transforms of x and y , then

$$r_{x,y} = \frac{\sum_{k=1}^n (x_k^A - \bar{x}^A) (y_k^A - \bar{y}^A)^*}{\left(\sum_{k=1}^n |x_k^A - \bar{x}^A|^2\right)^{1/2} \left(\sum_{k=1}^n |y_k^A - \bar{y}^A|^2\right)^{1/2}}, \quad (5)$$

where \bar{x}^A denotes the mean of x^A , $|\cdot|$ denotes absolute value, and $(*)$ denotes complex conjugation. Note that $r_{x,y}$ is complex-valued and that $|r_{x,y}| \leq 1$, where an absolute value of 0 and 1 correspond to uncorrelated and completely correlated, respectively. Also, its angle corresponds to the phase-delay between x and y . In particular, $r_{y,x} = r_{x,y}^*$, where $*$ denotes complex conjugation. The *absolute imaginary Pearson correlation coefficient* between x and y , denoted by $r_{x,y}^{\text{imag}}$, is obtained by taking the absolute value of the imaginary part of $r_{x,y}$:

$$r_{x,y}^{\text{imag}} = |\text{Im}(r_{x,y})|. \quad (6)$$

Similarly, the *absolute real Pearson correlation coefficient* between x and y , denoted by $r_{x,y}^{\text{real}}$, is obtained by taking the absolute value of the real part of $r_{x,y}$:

$$r_{x,y}^{\text{real}} = |\text{Re}(r_{x,y})|. \quad (7)$$

Note that $r_{x,y}^{\text{real}}$ and $r_{x,y}^{\text{imag}}$ measure the instantaneous and lagged functional connectivity respectively, and take values within the interval $[0, 1]$. By calculating the real and imaginary Pearson correlation coefficients for all pairs of cortical regions, we obtained n -dimensional cortical FC matrices $\text{RC}_{\text{cortex}}$ and $\text{IC}_{\text{cortex}}$, respectively, where n is the number of cortical regions. Furthermore, by calculating the real and imaginary Pearson correlation coefficients between the time-series of every pair of MEG channels, we obtained s -dimensional sensor FC matrices, denoted by $\text{RC}_{\text{sensor}}$ and $\text{IC}_{\text{sensor}}$, respectively, where s is the number of sensors. In fact, we obtained two sensor FC matrices, corresponding to axial and planar gradiometer data.

As a third measure, we take the *phase-lag index* (PLI), which is a measure of the asymmetry of the distribution of phase-differences between x and y and takes values in the interval $[0, 1]$. A value of 0 corresponds to no coupling or instantaneous coupling (that is, phase-difference centered around $0 \bmod \pi$), and a value of 1 corresponds to perfect phase locking at an angle $\neq 0 \bmod \pi$ [47]. Let $\psi = (\psi_1, \dots, \psi_n)$ denote the phase-differences between x_A and y_A . The phase-lag index between x and y is then defined as

$$\rho_{x,y} = |\langle \text{sign}(\psi_k) \rangle|, \quad (8)$$

215 where $\langle \cdot \rangle$ denotes the average over k . By calculating $\rho_{x,y}$ between the time-series of every pair of cortical
 216 regions or MEG sensors, we obtain FC matrices $\text{PLI}_{\text{cortex}}$ and $\text{PLI}_{\text{sensor}}$. As for the Pearson correlation
 217 coefficient, we calculated the sensor-level FC matrix for both the axial and planar gradiometer data.
 218 In the two following sections, we describe the methods we used for characterizing the FC matrices
 219 described above and for comparing corresponding cortical and sensor FC matrices.

220 2.4 Network analysis

221 We characterize and compare the topological features of the cortical and MEG sensor-level FC matri-
 222 ces, considering commonly used indices in MEG (and EEG) sensor-level network-theoretical studies as
 223 in e.g. [45].

224
 225 *Clustering coefficient:* The clustering coefficient is the likelihood that neighbors of a given node will be
 226 connected to each other [51]. This measure reflects the organization of the network: whereas a random
 227 network will have clustering coefficient equal or close to 0, one with a high degree of organization (e.g.
 228 a small-world network) will have a large clustering coefficient. Consider a matrix C whose $(i, j)^{\text{th}}$ entry
 229 c_{ij} specifies the correlation measure between point i and point j . The weighted clustering index of
 230 node i is defined as

$$c_i = \frac{\sum_{k \neq i} \sum_{\substack{l \neq i \\ l \neq k}} c_{ik} c_{il} c_{kl}}{\sum_{k \neq i} \sum_{\substack{l \neq i \\ l \neq k}} c_{ik} c_{il}} \quad (9)$$

231 whereas the mean clustering coefficient is given by

$$c_w = \frac{1}{N} \sum_{i=1}^N c_i. \quad (10)$$

232 To highlight the differences with respect to a random network, we compute the absolute difference
 233 $|c_w - \bar{c}_w|$, where the value \bar{c}_w is the clustering coefficient of a random network, which is equal to the
 234 network density.

235
 236 *Average path length:* this measure is the average number of steps along the shortest paths for all
 237 possible pairs of nodes in the network, and is a measure of the efficiency of information transport on
 238 the network [39]. The average path length depends on the network size but does not change drastically
 239 with it. We define the length of an edge as

$$\begin{aligned} l_{ij} &= \frac{1}{c_{ij}}, & \text{if } c_{ij} \neq 0, \\ l_{ij} &= \infty, & \text{if } c_{ij} = 0 \end{aligned}$$

240
 241 The average weighted path length is given by

$$l_w = \left(\frac{1}{N(N-1)} \sum_{i=1}^N \sum_{j \neq i}^N l(i, j)^{-1} \right)^{-1} \quad (11)$$

242 *Global efficiency:* efficiency is a measure of how efficiently the network exchanges information, it quan-
 243 tifies the exchange of information across the network. The global efficiency is computed as the average
 244 of the inverse shortest path length [39].

245
 246 Communities are groups of nodes within a network that are more densely connected to one another than
 247 to other nodes. The metric modularity quantifies the quality of an assignment of nodes to communities
 248 by evaluating how much more densely connected the nodes within a community are compared to how

249 connected they would be, on average, in a random network. We consider therefore an analysis which
250 may shed additional light on the group organization of the network: the Louvain community detection.

251

252 *Louvain community detection:* The Louvain method is an algorithm for detecting communities in
253 networks that relies upon heuristics for maximizing the modularity [3]. The optimal community struc-
254 ture returned by the algorithm is a subdivision of the network into non-overlapping groups of nodes,
255 such that the number of within-group edges is maximized, whereas the number of between-group edges
256 is minimized. The size and the number of communities detected is controlled by the resolution pa-
257 rameter, with standard value equal to 1. Whereas resolutions higher than 1 produce larger number of
258 clusters, resolutions lower than 1 produce lower number of clusters.

259 3 Results

260 3.1 Simulated resting-state cortical activity

261 We simulate resting-state cortical oscillations in the alpha frequency band (≈ 10 Hz) for a total of five
262 minutes using the Hopf oscillators coupled through the structural connectivity matrix S , obtained from
263 diffusion MRI fiber-tracking (see Section 2.1). We want to investigate whether there is a correlation
264 between the FC matrices at source level and the FC matrices at sensor level; therefore, to fit the 273
265 sensors of the MEG helmet we choose a brain parcellation composed of $n = 219$ ROIs, listed in Table 1.
266 Fig. 2 (a) shows the empirical structural connectivity matrix S , which exhibits a sparse connectivity,
267 whereas Fig. 2 (b) shows the instantaneous FC matrix, denoted RC_{cortex} . Observe that the first
268 quadrant of the matrix S contains the connectivity information from the right hemisphere, whereas
269 the fourth contains the connectivity information from the left hemisphere. Moreover, the second and
270 third quadrants of the matrix show the connections between the left and right hemispheres. The brain
271 model described in Sec. 2.1 is implemented with the parameter values listed in Table 2, and a time
272 sample of brain activity is shown in Fig. 2 (c) where different colors are used to distinguish between
273 nodes with different activity. A seed-based RC_{cortex} matrix has been computed for the right Caudal
274 Middle Frontal 1 region and mapped onto the different brain areas, as shown in Fig. 2 (d).

275 Fig. 3 (a) depicts the time series generated for 10 seconds of brain activity at one node of the brain
276 model, and Fig. 3 (b) shows the MEG signal obtained with axial gradiometer sensors for consecutive
277 time samples as measured at sensor level (plots are read from left to right and top to bottom). The
278 propagation patterns observed in Fig. 3 (b) suggest a spatial organization of the relative latencies in
279 the oscillations recorded at the different sensors. From these images we may infer that a more natural
280 way to conceptualize sensor-level resting-state MEG brain dynamics is as a spatiotemporal continuum
281 sampled in space and time by the channels [24]. Motivated by these observations, we investigate if a
282 network-theoretical approach might be appropriate to characterize them.

283 3.2 Comparison of cortical and MEG functional networks

284 In this section we systematically compare the network-theoretical properties of the simulated FC ma-
285 trices at the cortical- and sensor-level using the network indices described in Sec. 2.4. As outlined in
286 Section 2.3, we use FC matrices computed using three different measures: instantaneous/real correla-
287 tion (RC), lagged/imaginary correlation (IC), and the phase-lag index (PLI). Moreover, sensor-level
288 activity is computed assuming either axial or planar gradiometer MEG sensors. After considering
289 some general properties of the cortical and sensor functional networks in Sec. 3.2.1, we characterize
290 and compare their clustering coefficients and global efficiency in Sections 3.2.2 and 3.2.3, respectively.
291 Finally, in Sec. 3.2.4 we compare their community structure.

RIGHT hemisphere		RIGHT hemisphere		LEFT hemisphere		LEFT hemisphere	
1	lateralorbitofrontal 1	57	superiorparietal 4	109	lateralorbitofrontal 1	165	superiorparietal 1
2	lateralorbitofrontal 2	58	superiorparietal 5	110	lateralorbitofrontal 2	166	superiorparietal 2
3	lateralorbitofrontal 3	59	superiorparietal 6	111	lateralorbitofrontal 3	167	superiorparietal 3
4	lateralorbitofrontal 4	60	superiorparietal 7	112	lateralorbitofrontal 4	168	superiorparietal 4
5	parsorbitalis 1	61	inferiorparietal 1	113	parsorbitalis 1	169	superiorparietal 5
6	frontalpole 1	62	inferiorparietal 2	114	frontalpole 1	170	superiorparietal 6
7	medialorbitofrontal 1	63	inferiorparietal 3	115	medialorbitofrontal 1	171	superiorparietal 7
8	medialorbitofrontal 2	64	inferiorparietal 4	116	medialorbitofrontal 2	172	inferiorparietal 1
9	medialorbitofrontal 3	65	inferiorparietal 5	117	parstriangularis 1	173	inferiorparietal 2
10	parstriangularis 1	66	inferiorparietal 6	118	parsopercularis 1	174	inferiorparietal 3
11	parstriangularis 2	67	precuneus 1	119	parsopercularis 2	175	inferiorparietal 4
12	parsopercularis 1	68	precuneus 2	120	rostralmiddlefrontal 1	176	inferiorparietal 5
13	parsopercularis 2	69	precuneus 3	121	rostralmiddlefrontal 2	177	precuneus 1
14	rostralmiddlefrontal 1	70	precuneus 4	122	rostralmiddlefrontal 3	178	precuneus 2
15	rostralmiddlefrontal 2	71	precuneus 5	123	rostralmiddlefrontal 4	179	precuneus 3
16	rostralmiddlefrontal 3	72	cuneus 1	124	rostralmiddlefrontal 5	180	precuneus 4
17	rostralmiddlefrontal 4	73	cuneus 2	125	rostralmiddlefrontal 6	181	precuneus 5
18	rostralmiddlefrontal 5	74	pericalcarine 1	126	superiorfrontal 1	182	cuneus 1
19	rostralmiddlefrontal 6	75	pericalcarine 2	127	superiorfrontal 2	183	pericalcarine 1
20	superiorfrontal 1	76	lateraloccipital 1	128	superiorfrontal 3	184	lateraloccipital 1
21	superiorfrontal 2	77	lateraloccipital 2	129	superiorfrontal 4	185	lateraloccipital 2
22	superiorfrontal 3	78	lateraloccipital 3	130	superiorfrontal 5	186	lateraloccipital 3
23	superiorfrontal 4	79	lateraloccipital 4	131	superiorfrontal 6	187	lateraloccipital 4
24	superiorfrontal 5	80	lateraloccipital 5	132	superiorfrontal 7	188	lateraloccipital 5
25	superiorfrontal 6	81	lingual 1	133	superiorfrontal 8	189	lingual 1
26	superiorfrontal 7	82	lingual 2	134	superiorfrontal 9	190	lingual 2
27	superiorfrontal 8	83	lingual 3	135	caudalmiddlefrontal 1	191	lingual 3
28	caudalmiddlefrontal 1	84	fusiform 1	136	caudalmiddlefrontal 2	192	lingual 4
29	caudalmiddlefrontal 2	85	fusiform 2	137	caudalmiddlefrontal 3	193	fusiform 1
30	caudalmiddlefrontal 3	86	fusiform 3	138	precentral 1	194	fusiform 2
31	precentral 1	87	fusiform 4	139	precentral 2	195	fusiform 3
32	precentral 2	88	parahippocampal 1	140	precentral 3	196	fusiform 4
33	precentral 3	89	entorhinal 1	141	precentral 4	197	parahippocampal 1
34	precentral 4	90	temporalpole 1	142	precentral 5	198	entorhinal 1
35	precentral 5	91	inferiortemporal 1	143	precentral 6	199	temporalpole 1
36	precentral 6	92	inferiortemporal 2	144	precentral 7	200	inferiortemporal 1
37	paracentral 1	93	inferiortemporal 3	145	precentral 8	201	inferiortemporal 2
38	paracentral 2	94	inferiortemporal 4	146	paracentral 1	202	inferiortemporal 3
39	paracentral 3	95	middletemporal 1	147	paracentral 2	203	inferiortemporal 4
40	rostralanteriorcingulate 1	96	middletemporal 2	148	rostralanteriorcingulate 1	204	middletemporal 1
41	caudalanteriorcingulate 1	97	middletemporal 3	149	caudalanteriorcingulate 1	205	middletemporal 2
42	posteriorcingulate 1	98	middletemporal 4	150	posteriorcingulate 1	206	middletemporal 3
43	posteriorcingulate 2	99	bankssts 1	151	posteriorcingulate 2	207	middletemporal 4
44	isthmuscingulate 1	100	superiortemporal 1	152	isthmuscingulate 1	208	bankssts 1
45	postcentral 1	101	superiortemporal 2	153	postcentral 1	209	bankssts 2
46	postcentral 2	102	superiortemporal 3	154	postcentral 2	210	superiortemporal 1
47	postcentral 3	103	superiortemporal 4	155	postcentral 3	211	superiortemporal 2
48	postcentral 4	104	superiortemporal 5	156	postcentral 4	212	superiortemporal 3
49	postcentral 5	105	transversetemporal 1	157	postcentral 5	213	superiortemporal 4
50	supramarginal 1	106	insula 1	158	postcentral 6	214	superiortemporal 5
51	supramarginal 2	107	insula 2	159	postcentral 7	215	transversetemporal 1
52	supramarginal 3	108	insula 3	160	supramarginal 1	216	insula 1
53	supramarginal 4			161	supramarginal 2	217	insula 2
54	superiorparietal 1			162	supramarginal 3	218	insula 3
55	superiorparietal 2			163	supramarginal 4	219	insula 4
56	superiorparietal 3			164	supramarginal 5		

Table 1: Brain parcellation composed of 219 ROIs.

Parameter	symbol	value
Number of areas	n	219
Bifurcation parameter	a	10
Global coupling constant	g	50-100
Areal oscillation frequency	f_0	10 ± 1 Hz
Areal noise intensity	σ	1×10^{-3}
sampling frequency	f_s	100 Hz
time	t	300 s
peak SNR of MEG data	PSNR	0 – 30 dB

Table 2: Parameters of the cortical Hopf model, their symbols, and nominal values.

3.2.1 General properties

Fig. 4(a)-(c) show the cortical FC matrices RC_{cortex} , IC_{cortex} , PLI_{cortex} , and the ensuing sensor FC matrices RC_{sensor} , IC_{sensor} , PLI_{sensor} , for both axial (Fig. 4(d)-(f)) and planar (Fig. 4(g)-(i)) gradiometer MEG sensors. Moreover, Fig. 4(j)-(l) depict the distributions of FC values for each of the three FC types. We can make a number of observations. First, notice that the RC cortical FC exhibits a similar structure than IC and PLI cortico FC, but significantly higher values. Furthermore, this is approximately true for the sensor activity as well, which agrees with experimental observations [33]. It means that latencies between oscillations at different areas or sensors are much smaller than the oscillation period (which is about 100 ms for alpha oscillations). Lagged cortical FC (IC and PLI) values, however, are larger than lagged sensor FC values, which means that sensor FC matrices underestimate lagged FC. The main cause of this reduction in lagged FC is the instantaneous mixing of cortical signals through field spread. This phenomena is not unique to MEG recordings but can be observed in EEG and local field potential (LFP) recordings as well [25]. Second, lagged functional networks as measured with PLI are sparser than as measured with IC, and the distribution of their values is less spread for the planar gradiometer system.

Concerning the gradiometer type, notice that the instantaneous functional network (RC) is sparser for planar gradiometers (Fig. 4(g)) than for axial gradiometers (Fig. 4(d)). This is to be expected since planar gradiometers yield the highest signal intensities right above active cortical tissue and integrate source activity over smaller regions of cortex [21]. We observe that the matrices at sensor level (Fig. 4 (d)-(i)) show more modularity and a higher level of organization, and as expected, from the first column (Fig. 4(a), (d), (g)) we can see that the real correlation matrix RC is more sensitive to field spread.

3.2.2 Topology of resting-state networks I: MEG axial gradiometers

The network indices computed from the FC matrices at source level are compared here with those obtained from the FC matrices at sensor level assuming the MEG system with axial gradiometers. In order to do so, we apply a threshold and compute the binary FC matrices to analyze their organization structure for different threshold values. Fig. 5 (a) shows the clustering coefficient for RC computed with (9) as a function of the binarization threshold, along with the density. Observe that for values in the range $[0.2, 0.9]$ the clustering coefficient of RC_{sensor} exhibits a sort of flooring effect, showing that, although the total number of connections decreases, the number of 3-dimensional cliques remains almost unchanged within that range. This is equivalent to say that the links connecting three nodes are robust. However, this effect is not observed in the RC_{cortex} where the clustering coefficient curve shows a decreasing trend more similar to the one for the density at sensor level. Fig. 5 (b) shows the absolute difference $|c_w - \bar{c}_w|$ for both RC_{cortex} and RC_{sensor} , where \bar{c}_w denotes the average clustering coefficient of a random network, which in this case is equal to the density. In other words, this measure quantifies the difference in clustering with respect to a random network. The larger the value obtained

329 is, the higher the organization of the network (for that particular threshold value). We identify (and
330 highlight) an *optimum* value for each curve, which clearly do not coincide at source and at sensor level.
331 Fig. 5 (c) shows the efficiency of the network as a function of the binarization threshold. Again, the
332 efficiency computed for RC_{cortex} evidences a different organization than the one for RC_{sensor} . From
333 these results we can infer that RC_{sensor} exhibits a higher level of organization than RC_{cortex} for values
334 above 0.35, or equivalently, less randomness. Different results are obtained for IC and PLI matrices,
335 where both IC_{cortex} and PLI_{cortex} exhibit larger clustering coefficients than those at source level in
336 IC_{sensor} and PLI_{sensor} . Note that the FC matrix exhibiting largest distance with respect to a random
337 network in this case is PLI_{sensor} followed by PLI_{cortex} , as shown in Fig. 5 (e). On the other hand, the
338 global efficiency for these matrices is highest for IC_{cortex} (see Fig. 5 (f)), followed by PLI_{cortex} .

339
340 The results presented so far are obtained considering noise-free measurements. To assess the impact
341 of measurement noise, we perform simulations for several values of PSNR, as defined in (4). Fig. 6
342 plots the results vs. PSNR values in the range [0, 30] dB: (a) and (b) show respectively the clustering
343 coefficient and the global efficiency vs. PSNR for the real correlation matrix RC_{sensor} , along with
344 the results for its randomized counterparts, obtained by randomly shuffling the entries of the upper
345 triangular part of the matrix. Fig. 6 (c) and (d) show the same results for IC and PLI. As expected,
346 the clustering coefficient is slightly larger for the FC matrices than their random versions in all cases,
347 while the global efficiency is slightly higher for the random versions in all the PSNR spectrum. In all
348 cases, we observe that the network indices increase with high PSNR, with a clear transition starting
349 from 10 dB. These incremental behaviors are also observed in the number of communities detected, as
350 explained in Sec. 3.2.4.

351 3.2.3 Topology of resting-state networks II: MEG planar gradiometers

352 In this section we compare the network indices assuming the MEG system with planar gradiometers.
353 Fig. 7 (a) shows the clustering coefficient for RC_{cortex} and RC_{sensor} as functions of the binarization
354 threshold. For the sake of clarity, we include again the curve for the density. Moreover, Fig. 7 (b)
355 shows the absolute difference $|c_w - \bar{c}_w|$ computed for both RC_{cortex} and RC_{sensor} . As expected, the
356 results obtained for the RC matrices are similar to those obtained with the axial gradiometer sensors,
357 whereas the results for IC and PLI matrices are slightly different. In Fig. 7 (e) we can see that in
358 this case, IC_{cortex} and PLI_{cortex} show less distance with respect to a random network than IC_{sensor}
359 and PLI_{sensor} . Regarding noisy data, Fig. 8 (a) and (b) show the clustering coefficient and the global
360 efficiency vs. PSNR in the range [0, 30] for the real correlation matrices RC, along with the results
361 for their randomized counterparts. Fig. 8 (c) and (d) show the same results for IC and PLI at both
362 source and sensor level. The clustering coefficient and the efficiency increase with PSNR for all FC
363 matrices. From Fig. 8 (c) and (d) we can see that both the clustering coefficient and the global
364 efficiency are higher for IC_{sensor} than for PLI_{sensor} . From all the results presented, we can infer that
365 the FC matrices at sensor level show a different level of organization than the FC matrices at source
366 level, independently of the gradiometer system used.

367 3.2.4 Community structure

368 The Louvain modularity analysis performed for all FC matrices and assuming different values of the
369 resolution parameter yields more interesting results. Table 3 lists the number of detected communities
370 for each network assuming the standard value of the resolution parameter (i.e., 1), and Fig. 9 provides
371 a graphical representation of these results, where the nodes belonging to the same community are
372 rearranged and the communities are reordered by size within the respective matrix. Note that a large
373 number of communities at the cortex are composed of only a few elements, whereas at sensor level,
374 the planar matrices have a smaller number of communities (last row of Fig. 9). Moreover, the axial
375 IC_{sensor} and PLI_{sensor} have a large number of communities with only one element. Fig. 10 shows the
376 results of mapping the Louvain communities detected onto the MEG helmet, where only the most six

$RC_{\text{cortex}} = 28$	$IC_{\text{cortex}} = 18$	$PLI_{\text{cortex}} = 20$
$RC_{\text{sensor}} \text{ axial} = 6$	$IC_{\text{sensor}} \text{ axial} = 51$	$PLI_{\text{sensor}} \text{ axial} = 28$
$RC_{\text{sensor}} \text{ planar} = 4$	$IC_{\text{sensor}} \text{ planar} = 14$	$PLI_{\text{sensor}} \text{ planar} = 8$

Table 3: Number of communities detected for each FC matrix.

377 significant communities are highlighted in colors (the remaining communities are all colored in black).
378 The first row of Fig. 10 depicts the distribution of communities for RC_{sensor} , IC_{sensor} and PLI_{sensor}
379 for the MEG system with axial gradiometer sensors, whereas the second row shows the distribution of
380 communities for RC_{sensor} , IC_{sensor} and PLI_{sensor} assuming the MEG system with planar gradiometer
381 sensors. Except for the case of RC (first column of Fig. 10), no particular patterns can be identified
382 for the imaginary correlation matrices, and the communities seem to follow a radial distribution rather
383 than a well-defined network. These images reveal that we should be careful in order not to draw wrong
384 conclusions about the underlying structure of the brain.

385

386 We investigate next the impact of the resolution parameter of the Louvain method on the number of
387 detected communities for all matrices. Fig. 11 (a) depicts the number of communities detected as a
388 function of the resolution parameter in the range $[0.5, 1.5]$ for all FC matrices with the axial gradiome-
389 ter system, where the results have been obtained averaging 500 independent realizations. Analogously,
390 Fig. 11 (b) shows the results obtained with the planar gradiometer system. As expected, the number
391 of communities detected increases with increasing resolution parameter in both cases. Except for RC,
392 the number of communities detected at the cortex is smaller than the number of communities detected
393 at sensor level, and this difference is more evident for RC and PLI matrices in both MEG systems.
394 The difference at source and sensor level for RC remains almost constant for the entire resolution range
395 for the axial gradiometer system. This might be interpreted as the planar system having a greater
396 impact on the RC measures as the resolution parameter of the algorithm is increased, and less on the
397 IC measures.

398

399 From these results we can see that, increasing the resolution we systematically obtain a larger number
400 of communities for all matrices. We can also observe that the number of communities is larger at
401 sensor level than at source level for lagged correlations, as shown by the curves. The implications of
402 the results in Fig. 11 (a)-(b) are: first, we observe a discrepancy between the community structure of
403 cortical resting-state activity and that of the ensuing sensor level measurements (namely a different
404 number of communities). Second, it shows that this discrepancy is systematic, that is, at source level
405 the number of communities is always larger than at sensor level. Third, this observation holds for
406 lagged FC matrices (IC and PLI) and different gradiometer systems (axial/planar).

407

408 Analyzing the impact of noise in the number of detected communities, we observe from Fig. 11
409 (c)-(d) that noise has greater impact on the IC_{sensor} , as more communities are detected with higher
410 PSNR for both systems. The impact of noise is lower for RC_{cortex} , where the number of communities
411 detected does not change significantly with PSNR.

412 4 Discussion

413 A common way to conceptualize and analyze sensor-level EEG and MEG resting-state data is in terms
414 of functional networks [1, 4, 19, 29, 38, 43, 45, 46]. In this approach, the nodes of the network correspond
415 to EEG or MEG sensors and (the strength of its) links correspond to the estimated values of an inter-
416 action measure between the time-series recorded at the corresponding nodes. The functional network
417 is subsequently analyzed using graph-theoretical indices [8, 23, 39]. Although the network-theoretical
418 approach to sensor-level EEG/MEG data is potentially interesting since resting-state cortical dynamics
419 is known to be comprised of a number of functional networks [1, 2, 46], it is not without methodolog-

420 ical challenges [28, 50]. One of these challenges, which has received surprisingly little attention, is
421 the fact that EEG/MEG record linear superpositions of cortical activity [21]. Although this nature
422 of EEG/MEG signals has motivated the development of interaction measures that are less affected by
423 volume-conduction/field spread [7, 26, 32, 33, 47], as well as the use of source reconstruction using tech-
424 niques such as beamforming [10, 18, 41, 49], surprisingly few studies have investigated the relationship
425 between cortical and sensor-level EEG/MEG networks [22, 41], despite the fact that these topologi-
426 cal features are of high interest in experimental studies [4–6, 15, 16, 19, 29, 45, 46]. Moreover, simple
427 simulations that consider only a small number of regions do not allow to assess network properties.
428 In this study we used a whole-brain computational modeling approach to resting-state dynamics [13]
429 to compare cortical and sensor-level resting-state MEG networks, and considering two different MEG
430 systems: sensors with axial gradiometers and sensors with planar gradiometers. In our simulations,
431 we included both instantaneous interaction, as measured by real-valued correlation, as well as lagged
432 interaction, as measured by imaginary-valued correlation and the phase lag index (PLI) [47]. In ad-
433 dition to the commonly studied network measures, we performed a Louvain community analysis and
434 we studied the impact of noise in the measurements using empty-room MEG recordings. Below, we
435 discuss our main findings and their implications for network-based analysis of MEG sensor-level data.

436

437 One of our main findings is that the effect of field spread on network topology depends on the type
438 of interaction (instantaneous or lagged), measured by different interaction measures. This holds both
439 for the magnitude of the effect as well as for how field spread changes network topology. In particu-
440 lar, instantaneous correlations are more susceptible for field spread than lagged correlations, which is
441 not surprising because field spread is instantaneous [21]. In fact, the interaction measures that have
442 been proposed to deal with volume-conduction or field spread [26, 32, 33, 47] all are based on lagged
443 correlations, in one way or another. But field spread also affects instantaneous and lagged interac-
444 tion in different ways: for instantaneous interaction, it leads to a "flooring" of the mean clustering
445 coefficient and global efficiency (as a function of binarization threshold) and to an increased threshold
446 for maximal distance to a random network. For lagged interaction, the mean clustering coefficient
447 and global efficiency decrease (for any given threshold). Thus, our simulations suggest that lagged
448 interactions are more easily assessed from sensor-level MEG data, a finding that generalizes previous
449 studies [26, 32, 33, 47] to large-scale functional networks.

450

451 The Louvain community analysis yields interesting results. First of all, at source level there is a
452 large number of modules with few elements, a particularity not observed at sensor level. For imagi-
453 nary correlation (IC) and phase-lag index (PLI) at source level there is only one big community. These
454 organizations do not reflect the coupling among nodes due to the structural connectivity matrix, but
455 rather reflect the spatial low-pass properties of the MEG forward model. The same effect is captured
456 by the axial system at sensor level. On the other hand, a significantly smaller number of communities
457 is detected by the algorithm for the planar system, which may lead to the conclusion of a more orga-
458 nized underlying structure. In general, the number of communities at source and sensor level do not
459 match, and the results suggest a different structural organization of the functional connectivity (FC)
460 matrices. Similar conclusions can be drawn from the case of noisy MEG measurements: as the peak
461 signal-to-noise (PSNR) increases, the number of communities increases and the results approach the
462 noise-free case. Regarding the network indices studied, their values are also affected by noise, showing
463 a discrepancy with respect to the noise-free case.

464

465 Another main finding is that the discrepancy between (sensor-level) MEG and cortical networks is
466 larger when using axial gradiometers than when using planar gradiometers. This holds for both in-
467 stantaneous and lagged interaction and for all studied network-theoretical measures (mean clustering
468 coefficient, network density, average path-length, and global efficiency). Most network-theoretical
469 studies using sensor-level MEG, however, have used axial gradiometers [1, 5, 17, 45], and (as far as we
470 know), (virtual) planar gradiometers have not yet been used for this purpose. To understand why
471 planar gradiometer data is less affected by field spread, recall that the (virtual) planar transformation

472 is the sum of the squared magnetic fluxes through two perpendicular directions that are (approx-
473 imately) perpendicular to the skull. This transformation, which is called the Laplacian, behaves like a
474 spatial high-pass filter. It hence reduces the MEG sensor’s sensitivity to distant sources because the
475 source’s magnetic fields contains predominantly low spatial frequencies. In contrast to MEG studies,
476 the Laplacian has been used frequently applied to (scalp) EEG data [48] as well as to local field po-
477 tential recordings [25, 31] to increase spatial resolution. Based on our findings, we recommend using
478 (virtual) planar gradiometer data when assessing network properties from sensor-level MEG data.

479 5 Conclusions

480 Functional networks constitute a common way to conceptualize and analyze sensor-level EEG and
481 MEG resting-state data, in which sensors are identified with nodes and the interaction between chan-
482 nel time-series with the network connections. The network-theoretical approach is however challenged
483 by the fact that EEG and MEG time-series are mixtures of source activity. Since still many contri-
484 butions found in the literature perform network analysis at sensor level, in this study we address the
485 question of to what extent the network topology can be reconstructed from sensor-level FC measures
486 in case of MEG data. Using a diffusion MRI-constrained whole-brain computational model of resting-
487 state cortical activity, we report discrepancies between source- and sensor-level network topologies,
488 and observe that the effect of field spread on network topology depends on the type of interaction
489 (instantaneous or lagged). Moreover, we found that instantaneous interaction is more sensitive to field
490 spread than lagged interaction, and that the discrepancies are reduced when using planar gradiometers
491 rather than axial gradiometers. We therefore recommend to use lagged interaction measures on planar
492 gradiometer data when investigating network properties of resting-state sensor-level MEG data.

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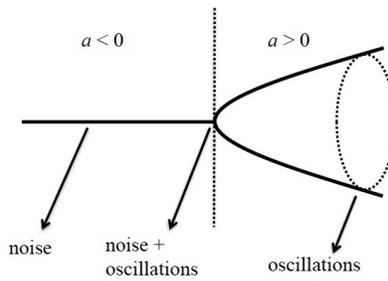
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(a) Bifurcation parameter

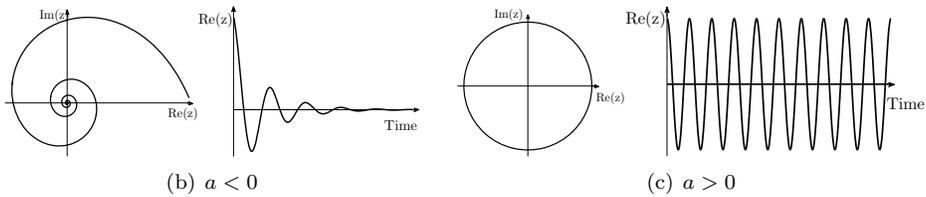


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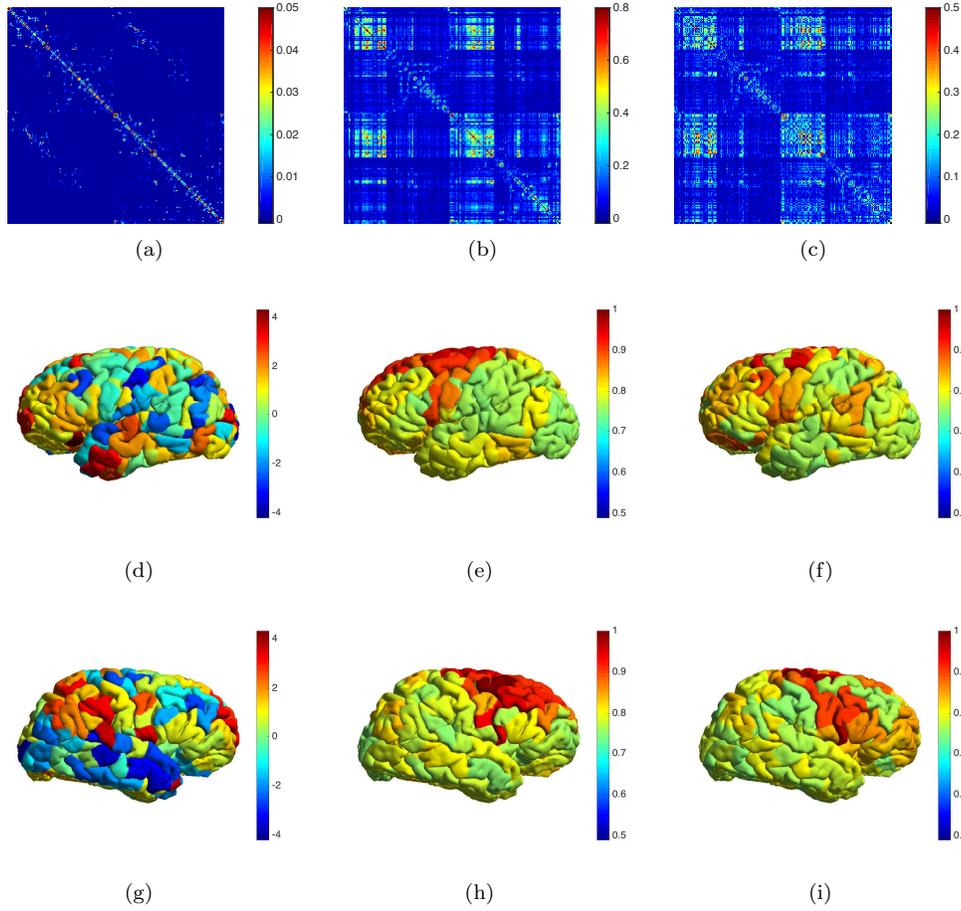


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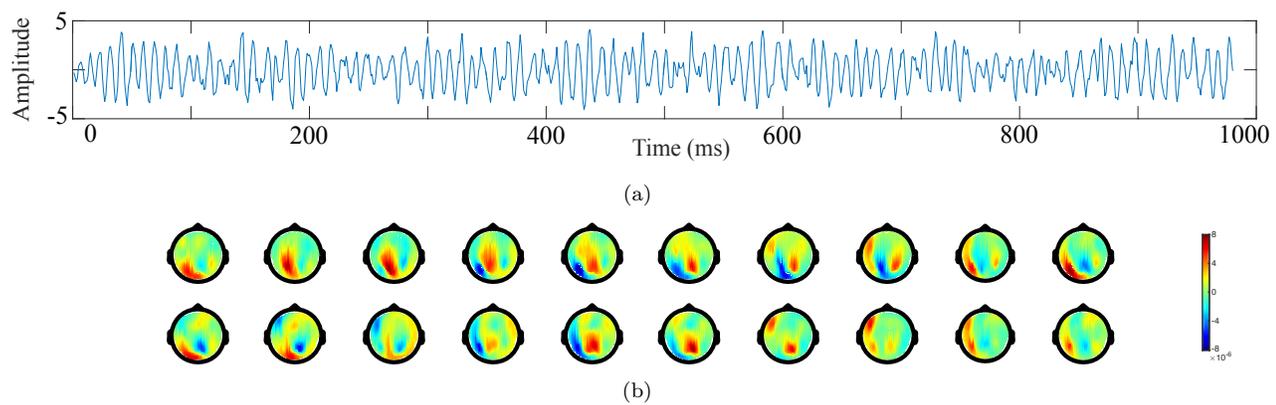


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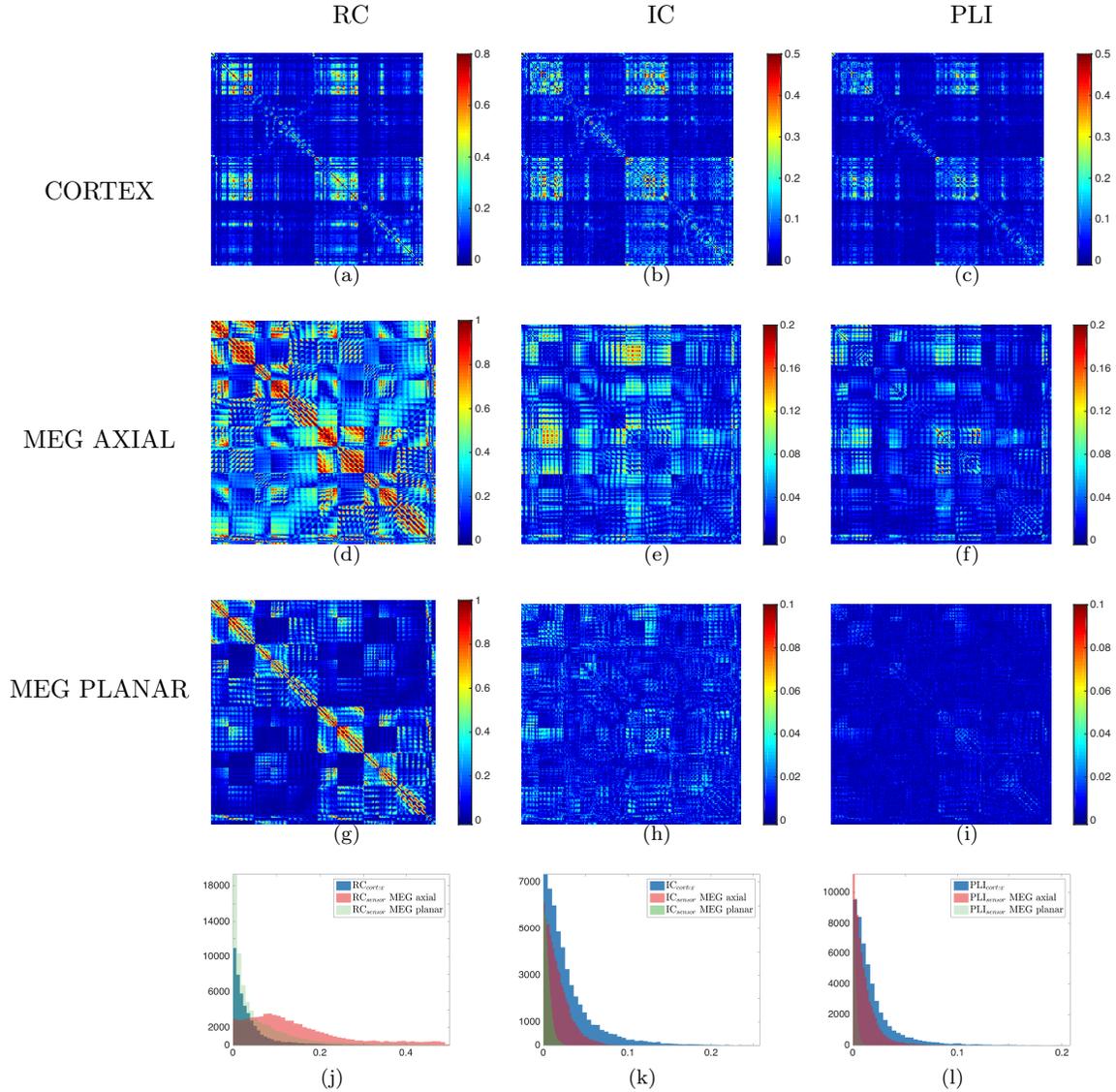


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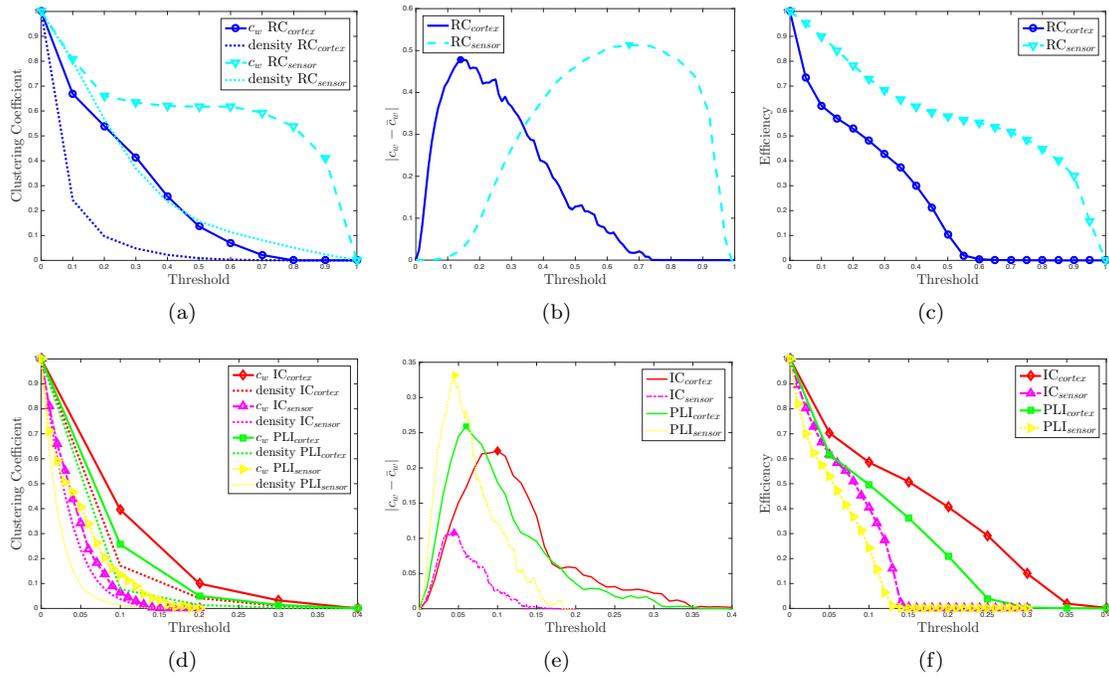


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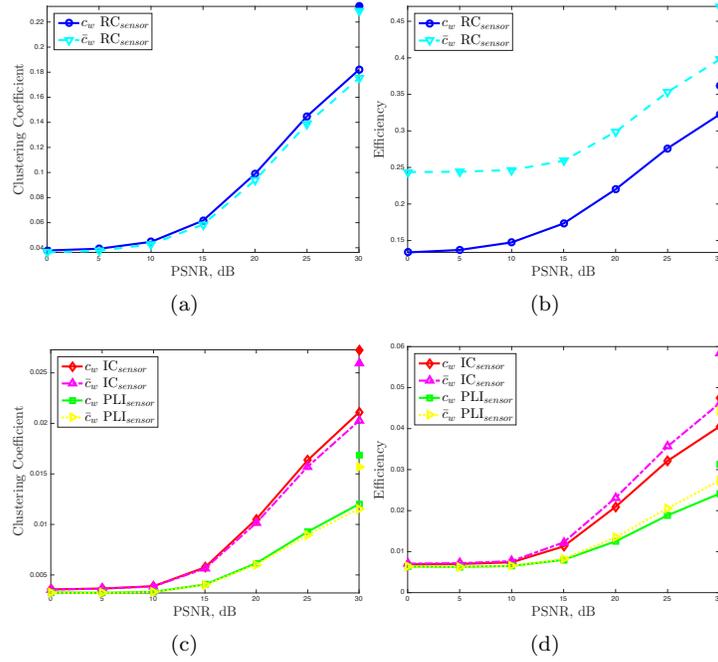


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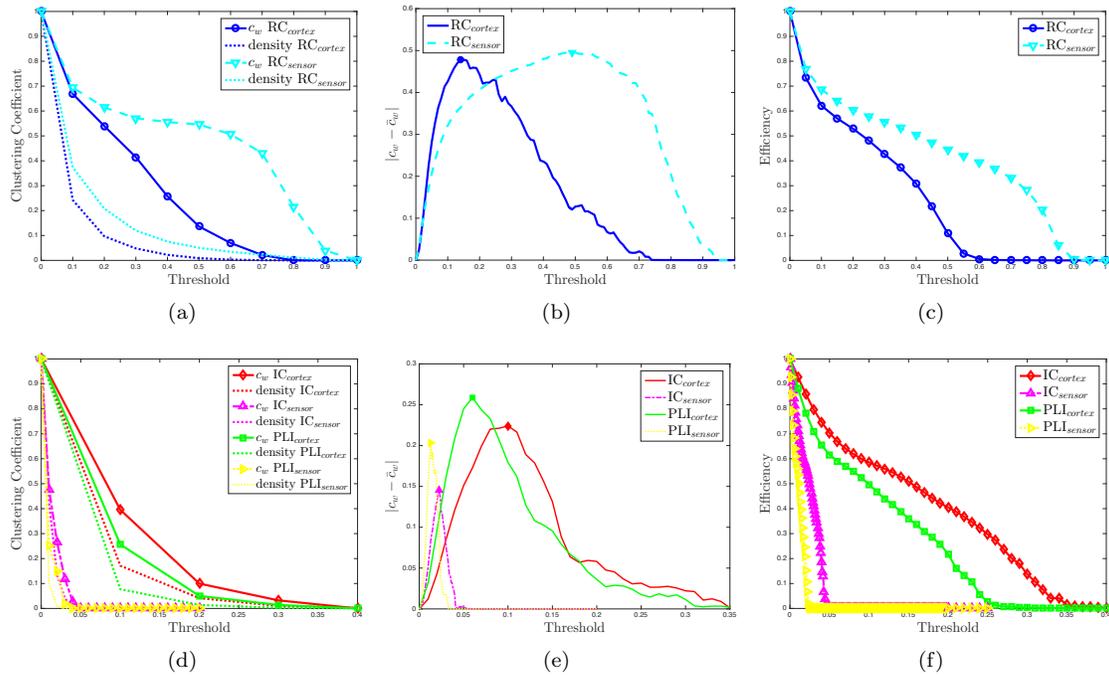


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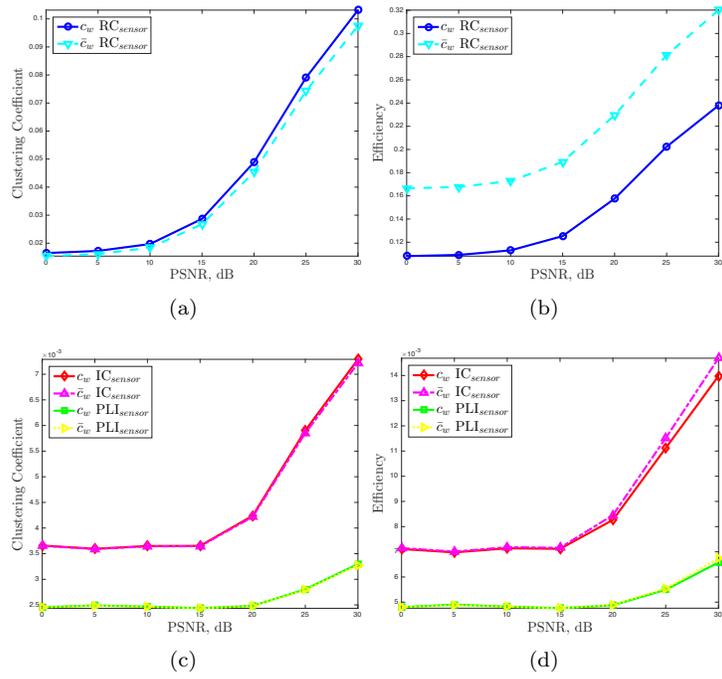


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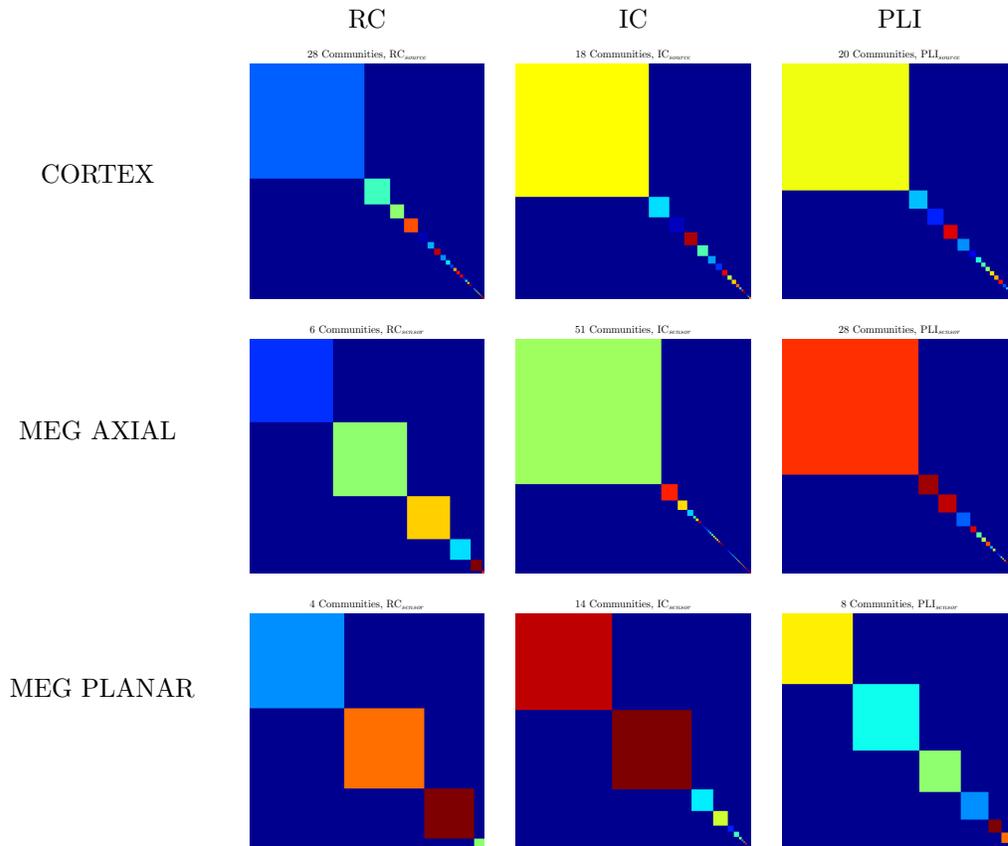


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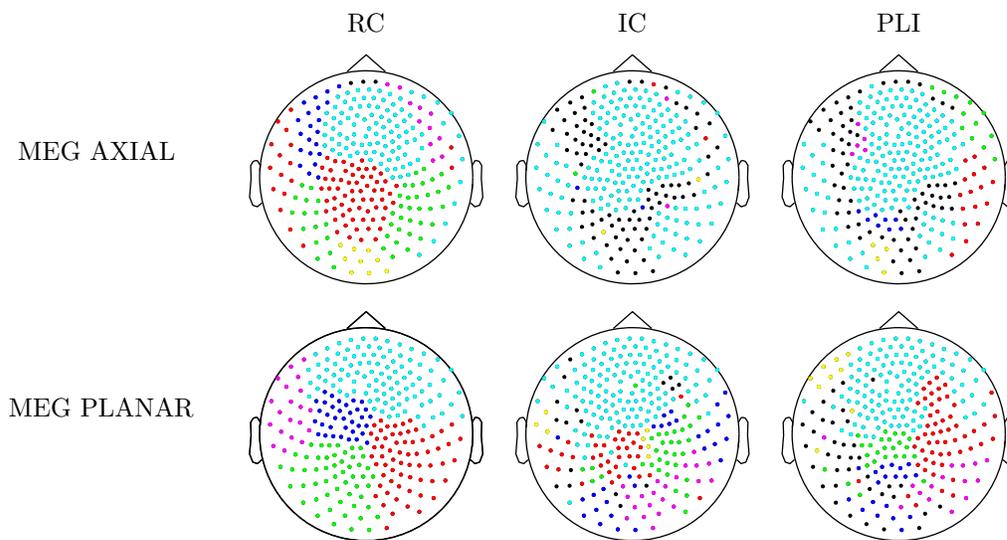


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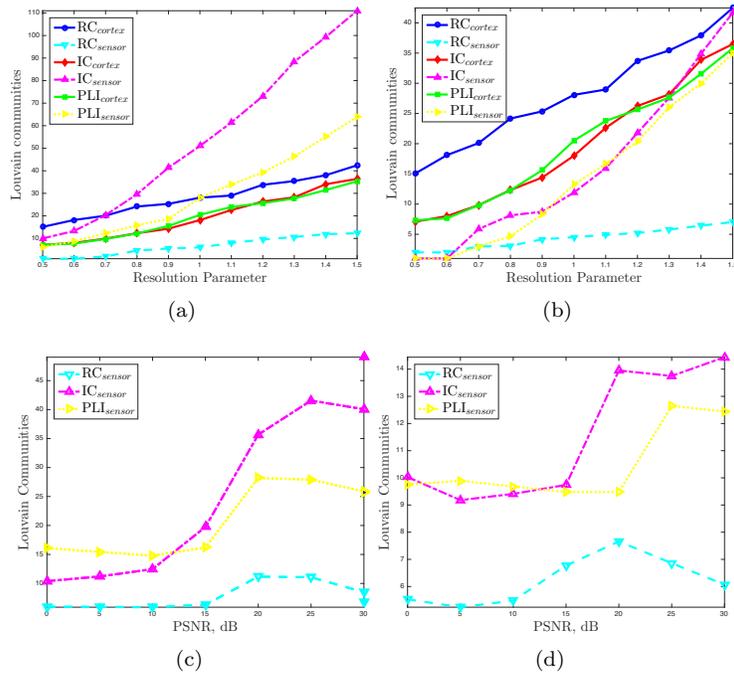


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