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Dimensionality Reduction Impedes the Extraction of Dynamic Functional Connectivity States from fMRI Recordings of Resting Wakefulness

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Abstract

Background—Resting wakefulness is not a unitary state, with evidence accumulating that spontaneous reorganization of brain activity can be assayed through functional magnetic resonance imaging (fMRI). The dynamics of correlated fMRI signals among functionally-related brain regions, termed dynamic functional connectivity (dFC), may represent nonstationarity arising from underlying neural processes. However, given the dimensionality and noise inherent in such recordings, seeming fluctuations in dFC could be due to sampling variability or artifacts.

New method—Here, we highlight key methodological considerations when evaluating dFC in resting-state fMRI data.

Comparison with Existing Method—In particular, we demonstrate how dimensionality reduction of fMRI data, a common practice often involving principal component analysis, may give rise to spurious dFC phenomenology due to its effect of decorrelating the underlying time-series.

Conclusion—We formalize a dFC assessment that avoids dimensionality reduction and use it to show the existence of at least two FC states in the resting-state.

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Author contributions statement

Acquisition of data: BP. Analysis and interpretation of data: MK, SC, BP. Drafting of the manuscript: MK, SC. MK, SC, and BP revised the manuscript critically for important intellectual content. All authors reviewed the manuscript.

Keywords

Resting-state functional magnetic resonance; dynamic functional connectivity (dFC); FC state analysis; spatiotemporal analysis

1. Introduction

Resting-state functional magnetic resonance imaging (fMRI) is a non-invasive tool to evaluate the large-scale organization of the brain when a subject is not performing an explicit task [1, 2]. Functional connectivity (FC) is a measure of association (usually, the correlation) between blood-oxygen-level dependent (BOLD) signals of distinct brain regions. Such analysis has revealed a number of resting-state networks (RSN) [3], each consisting of regions grouped by putative functional relevance as they recapitulate activation patterns of task-based paradigms.

Changes in resting-state FC have been reported at time scales of seconds to minutes [4, 5]. Thus, a sliding time window analysis has been increasingly used in the literature in order to study temporal fluctuations in FC, termed dynamic functional connectivity (dFC) [6, 7, 8, 9, 10, 11]. Within this framework, the obtained FC patterns are interpreted as a set of discrete states, called “FC states,” each representing a different configuration of RSNs during resting wakefulness [12, 13, 14, 15, 16, 17, 18, 19]. From this point of view, wakefulness is characterized by the sequential, but potentially stochastic, traversal of FC states [20].

However, the nature of fMRI recordings has complicated the development of dFC methodologies. For example, scanner drift, head motion, and physiological noise [21, 22], may lead to the appearance of signal nonstationarity [23, 24]. Moreover, it is unclear whether the temporal precision of the BOLD signal would be adequate to represent FC states in the brain. However, despite these potential confounds, it is argued that proper parameter choice (e.g., sliding window length) can enable detection of legitimate neurophysiological dFC temporal fluctuations [25, 26, 27].

Another potential issue regarding dFC methods is a statistical confound due to sampling variability [28]. In this scenario, data that are mathematically stationary and originate from a single FC state may give rise to multiple discrete FC states due to inconsistent sampling. Here, we examine this potential confound with specific regard for dimensionality reduction of the BOLD time-series. This processing step is often performed via principal components analysis (PCA) to reduce computational burden or to remove spurious artifact and noise [29, 30, 31, 32]. PCA linearly transforms the original fMRI data into a basis of mutually uncorrelated components [33]. Here, we consider temporal PCA, as utilized in [28], which reduces the spatial dimensionality in the transformed domain by retaining the components with high variability [34]. In other words, after applying temporal PCA to the BOLD signal, we obtain a signal in the transformed domain with the same number of time points and reduced number of regions in the spatial dimension. We will proceed to show that applying PCA in this way prior to FC state analysis can lead to interpretations that do not fully account for the structure of the data.

Specifically, our contributions are: (i) We illustrate that dimensionality reduction of the analyzed signals, due to its decorrelating action, may lead to inadequate assessment of dFC by promoting the separation of windows into clusters of differing statistics; (ii) We show that in the absence of dimensionality reduction, the composition (i.e. correlation matrix) *and* occupancy (i.e., percent of the time) of a FC state must be considered when testing for the validity of FC states; (iii) We show that incorporating these considerations allows for the extraction of at least two interpretable FC states during resting wakefulness.

2. Materials and Methods

2.1. Participants

This is a secondary analysis performed following an initial investigation on the average changes in functional connectivity following the transition between wakefulness and anesthetized states [35]. All aspects of this study were conducted under a protocol approved by the Washington University Human Research Protection Office. All the participants were between 18 to 40 years of age. The following exclusion criteria were considered in this study: use of over-the-counter antihistamines, antiemetics, herbal supplements, ethanol, or illicit drugs within a week of the study; disturbance in normal sleep pattern within the past 14 days; sedation, general anesthesia, or upper respiratory infection in the past 30 days; known auditory or physical impairments; known family history of malignant hyperthermia; implanted magnetic resonance imaging (MRI)-incompatible metal or prosthetic.

2.2. Experimental BOLD Data Acquisition and Pre-processing

We report analyses of fMRI data acquired from 10 human volunteers (Siemens 3T Trio, TR 2.2s, TE 27ms, flip angle = 90 degrees, 36 slices, 200 volumes/run). Two echo-planar BOLD imaging runs were acquired from each volunteer during quiet eyes-closed passivity (resting-state), prior to induction of general anesthesia. Pre-processing of data was performed as previously described [35]. Epochs contaminated by artifact related to micromovements of the head were censored. Regression of the whole brain global signal was also carried out. It should be noted that although global signal regression remains a controversial technique, it has been shown that it decreases the spurious changes in correlated BOLD signals that are introduced by head motion [36, 37, 38].

2.3. Cortical Segmentation into Resting-State Networks

In this study, $6 \times 6 \times 6$ mm regions were used to segment cortical gray matter. 1076 regions were selected using a winner-take-all algorithm with at least 50% gray matter and more than 90% probability of exclusive assignment to one of the seven RSNs. A previously published parcellation method [39] was used to assign brain regions to each RSN. A brief summary of this method is as follows: a multi-layer perceptron was trained to associate BOLD correlation maps corresponding to predefined seeds with specific RSN identities. In this study, we considered seven RSNs including the Default Mode Network (DMN), Dorsal Attention Network (DAN), Ventral Attention Network (VAN), Frontal Parietal Control Network (FPC), Visual Network(VIS), Language Network (LAN), and Somatomotor Network (SMN). DMN constitutes brain areas that show higher activity at rest than while performing a task [40, 41]. Maintaining attention to locations or features are thought to be

associated with DAN [42, 43]. VAN is considered to be involved in detecting novel stimuli and initiating shifts of attention [44, 45, 46]. FPC is hypothesized to be associated with working memory. Also, the FPC is thought to be responsible for configuring the brain's moment-to-moment information processing [47]. Visual and language processing are associated with VIS and LAN, respectively [48, 49, 50]. SMN is thought to be responsible for motor and sensory processing [45, 46, 51]. We then selected 15 representatives of each RSN by repeated pseudo-random re-sampling of regions from different RSNs. These sub-sampled data (105 voxels) have been made zero-mean and then used in our FC state analysis. In all of the figures presented in this paper, the assignment of 105 voxels to seven RSNs are as follows: DAN: 1–15; VAN: 16–30; SMN: 31–45; VIS: 46–60; FPC: 61–75; LAN: 76–90; DMN: 91–105.

2.4. Functional connectivity processing

In all analyses presented in this paper, the BOLD signals were band-pass filtered between 0.008 and 0.1 Hz using a 10th order Butterworth filter [52]. All the data analyses were performed using MATLAB (The MathWorks, Inc., MA, USA) scripts generated for this study.

2.5. Dynamic functional connectivity

In order to characterize the temporal dynamics of functional connectivity, we used a sliding window analysis of the BOLD signal [20, 21]. Computing correlations on shorter sub-trial windows (i.e., binning) can provide more temporal information at the expense of increased noise susceptibility [6]. We estimated the sliding window functional connectivity matrix for each time window. In this study, we used 50 samples (110 seconds) as the sliding window lengths to find FC states. For all the analyses, overlapping was performed with a separation of 5 samples (11 seconds) between consecutive sliding window centers.

2.6. FC state analysis of resting wakefulness

To evaluate FC states across subjects, we applied the k-means clustering technique to the correlation matrices obtained from all ten subjects. Thus, 737 sliding windows (with 110 seconds for the length of each window) from all subjects were used in the clustering. Correlation values between all region-pairs were included, resulting in $(105 \times 104)/2 = 5460$ features per matrix. The clustering algorithm was performed with random initialization of centroid points and was repeated 100 times to increase chances of escaping local minima. As the simplest scenario to assess altered FC states dynamics during resting wakefulness, we selected the number of clusters $k = 2$ in the k-means algorithm.

2.7. Simulated BOLD Data

We generated different cases of synthetic BOLD signals, described below, to generate positive and negative controls for the presence of FC states in the signals. To control for the spectral content, we matched the power spectrum of the synthetic data to that of the experimental BOLD signal.

- I. Simulated Stationary BOLD with corrected power spectrum. We generated stationary simulated BOLD fMRI time-series following [28]. Specifically, first

we generated random samples from a normal distribution. We used the same number of time points and voxels as in the experimental BOLD data for each subject. Then, we filtered these time-series utilizing the average power spectrum calculated from the experimental BOLD data. Then, we matched the simulated time-series covariance matrix to the one from the experimental data (by projecting the simulated time-series data onto the eigenvectors of the covariance matrix from the experimental data calculated over the entire scan time). By doing so, we generate a stationary synthesized time-series for all subjects in the dataset that has a static FC (time-averaged FC) and frequency content similar to the static FC and the average power spectrum of the actual BOLD signal, respectively. The static FC of each subject, calculated over the entire scan time, is provided in the Appendix (Figure 6).

- II. Simulated Non-stationary BOLD with corrected power spectrum. A non-stationary simulated BOLD fMRI time-series is generated based on the FC state analysis results on the experimental BOLD signal. For each subject, we consider two FC states with the assumption that the subject is transitioning between these two FC states over time during resting wakefulness (see the Appendix for more details). In this case, we extracted the FC states (and corresponding FC state assignment) for each window using the method described in section 2.6 for each individual subject. Then for each window, we generated a stationary time-series which match the covariance of the assigned FC state. Again we forced the power spectrum of the simulated BOLD to be similar to the average power spectrum of the actual BOLD signal. This is our positive control for the presence of FC states.
- III. Simulated Stationary BOLD with uncorrected power spectrum. We generated stationary simulated BOLD fMRI time-series for each subject as in Case I, but we did not modify the power spectrum of the simulated signal.

2.8. FC state occupancy, entropy and similarity

For a particular individual, we define the occupancy of a FC state as the ratio of windows assigned to that FC state relative to the total number of sliding windows. We denote the occupancy of subject i being at FC state j , $j = 1, 2$ as p_i^j , which is bounded between 0 and 1.

The entropy of occupancy for subject i is defined as $E_i = - \sum_{j=1}^2 p_i^j \log_2 p_i^j$. A zero entropy means that there are no transitions between FC states while a unit value for entropy means that the subject is revisiting all FC states with equal probability. We calculated the correlation between FC states for each case of simulated BOLD separately to compare FC states obtained from the experimental BOLD signals to FC states obtained from different simulated BOLD time-series. To quantify the relation between FC states similarity and entropy of occupancy in simulated BOLD cases I–II, we conducted a regression analysis to quantify the beta value (β), the R^2 and a p-value (P).

2.9. Statistical Analysis

We compare FC states obtained from the experimental BOLD signals to FC states obtained from each different simulated fMRI time-series.

Differences in occupancy of being at a particular FC state across different BOLD time-series, were assessed using a two-sample Kolmogorov–Smirnov test (K–S test). The two-sample K–S test is a nonparametric test for cases where the sample distributions are unknown which returns a test decision for the null hypothesis that two samples are from the same continuous distribution. Self-similarity of FC states across different BOLD time-series data, real and simulated, were tested using t-test on the z-values of FC states.

3. Results

3.1. Simulated and experimental BOLD signal can give rise to similar FC states

We first compare dFC state analysis on the experimental BOLD signal and the synthesized data from the different controls. For each of the positive and negative controls, we perform analysis with and without the application of PCA to the initial time-series data. When PCA is utilized, the original time-series data are dimensionality reduced from 105 to 30 dimensions prior to clustering, as in [28]. In our data, the percentage of variance that was retained after PCA reduction was 92.2% on average. We consider $k = 2$ for the k -means analysis.

Figure 1 illustrates the correlation matrices corresponding to the two FC states, FC state 1 (FCS1) and FC state 2 (FCS2), obtained for the simulated and experimental BOLD data. It can be concluded from this plot that the FC states from all controls are indeed similar to the ones obtained from the experimental BOLD signals. The pairwise similarity of FC states from the experimental BOLD signals to the simulated BOLD data from case I, case II and case III (with 100 realizations per case) was 0.89 ± 0.04 , 0.97 ± 0.01 , and 0.90 ± 0.02 (mean \pm SD), respectively. Thus, control cases can indeed generate FC states that appear similar to those obtained from the actual data. However, the difference between FC states is obscured following the use of PCA, which tends to homogenize the extracted FC states (Figure 2B). This plot suggests that FC states are less similar without PCA, while with the application of PCA, FC states manifest greater similarity.

3.2. FC state occupancy can differentiate simulated and experimental signals and is obscured by PCA

While the above is suggestive of a potential confound (i.e., stationary data giving rise to apparent FC states), to fully study the validity of FC states we must consider occupancy in addition to the FC state similarity. The FC state occupancy (i.e., the fraction of windows assigned to that FC state) across different subjects for the experimental BOLD signal, and control cases I–III are shown in Figure 2A.

It can be concluded from this plot that occupancy of either of the FC states in the experimental BOLD signal is entropic, and each subject exhibits transitions between these two FC states during the recording. The same entropic behavior is observed in the non-

stationary synthesized time-series (i.e., case II), as expected. However, for both case I and case III, the FC state occupancy obtained without using PCA shows mostly one FC state for each subject, which means that simulated subject does not exhibit any transitions between the two FC states. Statistical analysis confirmed that the occupancy of the first FC state is significantly different between the experimental BOLD signal and simulated stationary BOLD data regardless of whether the power spectral properties have been simulated in case I (K-S test, $P = 0.031$, $\text{stat} = 0.6$) or not as in case III (K-S test, $P = 0.007$, $\text{stat} = 0.7$). Moreover, the experimental BOLD signal is *not* significantly different from case II (K-S test, $P = 0.975$, $\text{stat} = 0.2$), where we simulate nonstationary BOLD data with power spectrum approximating that of the experimental signals.

However, application of PCA to the time-series data prior to FC state extraction changes the occupancy, especially for the stationary simulated signals (case I, III). We further quantified the FC state similarity (correlation between two FC states) and FC state entropy for cases with and without PCA, shown in Figure 2B,C. We tested the difference between FC states similarity with and without PCA by utilizing t-tests on z-values of correlations (applying Fisher inverse transform to correlation values) over subjects. The statistical results show significant difference with and without PCA for all the cases I–III with $P \ll 0.001$. We further analyzed the differences in the occupancy entropy over subjects, with and without applying PCA to the original time-series data. The results of such statistical analyses are as follows: (K-S test for the experimental BOLD, $P = 0.111$, $\text{stat} = 0.5$), (K-S test for case I, $P \ll 0.001$, $\text{stat} = 1$), (K-S test for case II, $P \ll 0.001$, $\text{stat} = 1$), (K-S test for case III, $P = 0.001$, $\text{stat} = 0.8$). The results of statistical tests on the occupancy, of being at a particular FC state, are reported in the Appendix. This analysis confirms that application of PCA does indeed homogenize FC states both in terms of their FC state similarities *and* in terms of their occurrences in time.

Figure 3A depicts the difference between two FC states, separately obtained with and without applying PCA to time-series data from the experimental BOLD signal and simulated BOLD in cases I–III. In Figure 3B, the FC state identity of each sliding window is projected onto the first two principal components. This figure clearly shows how applying PCA can fundamentally alter the distribution of time epochs to different FC states.

3.3. Accounting for occupancy reveals RSN-specific differences in FC states

The above methodological considerations indicate that when occupancy is fully taken into account, the experimental BOLD signal does indeed appear to exhibit transitioning between FC states. To evaluate the salience of these FC states from a physiological perspective, we performed a brief analysis to identify the key differences in the FCS1 and FCS2. We found that the most prevalent difference between these FC states manifests in the correlation between the DAN and VIS RSNs (Figure 4A). Specifically, we note that strong correlation between VIS and DAN in the first FC state is absent in the second one.

This connectivity difference between the first and second FC states for all the connections ($30 \times 29/2 = 435$ connections) are shown in Figure 4A for region-pairs within DAN and VIS networks, and was significant by paired t-test ($p \ll 0.001$) after correction for multiple comparisons ($30 \times 29/2 = 435$ connections). Figure 4B shows the comparison of connections

within DAN, connections within VIS, and connections between DAN and VIS. A two-way ANOVA (on z-values of correlations) with FC states (first or second FC state) and network (within DAN, within VIS, and between DAN and VIS) as factors shows a significant main effect of FC state ($F_{1,864} = 482.62, P \ll 0.001$), a significant main effect of network ($F_{2,864} = 659.38, P \ll 0.001$), and a significant network by FC state interaction ($F_{2,864} = 28.54, P \ll 0.001$).

3.4. Decorrelating actions of PCA lead to spurious FC state dynamics

The above analyses reveal a clearly problematic methodological step in FC state analysis, i.e., the application of dimensionality reduction prior to state extraction. The observed effects can potentially be understood in terms of the decorrelating actions of PCA. To substantiate this notion, we also performed analysis on a toy example consisting of non-stationary data randomly generated from two different noise-driven linear dynamical systems f_1 and f_2 (red and blue colors) (see the Appendix for the exact specification of f_1 and f_2). A random switch determines which dynamical system is sampled at each window. Sliding correlation analyses of the synthesized data without and with PCA are shown in Figure 5B and 5C, respectively. The correlation in 5C fluctuates about a mean of zero, as expected due to the effect of PCA. We then applied the FC state analysis and observed substantial errors in finding the true FC states after employing PCA (Figure 5D). Figure 5E shows the pairwise correlation for both FC states with and without PCA. The estimated correlation error is highlighted in Figure 5F. The exact covariance and correlation matrices, alongside the estimated ones, with and without PCA are reported in the Appendix.

4. Discussion

We evaluated the presence of FC states in the experimental BOLD signal acquired during eyes closed wakefulness and whether conventional cluster-based extraction methods could be susceptible to obscuring their assessment. We showed that dimensionality reduction (PCA) may be a problematic analysis step insofar as it could change the FC state similarity and occupancy of the extracted FC states. Taking this into account, we showed that FC states can, in fact, be reliably deduced in different positive controls of non-stationary simulated BOLD data and a toy example. Further, FC states extracted from experimental BOLD signals are interpretable from a physiological perspective.

4.1. Limitations of PCA in dFC analysis for FC state extraction

PCA is commonplace in FC analysis, given the high dimensionality of signals generated by BOLD fMRI and high-density EEG. It should be noted that similar to FC state extraction in BOLD fMRI data, EEG/MEG microstate analysis has been carried out for cases of high temporal resolution data [53, 54]. However, such a step has the effect of providing linearly uncorrelated signals throughout a recording. Thus, subsequent window-based clustering of the post-PCA data will, almost by definition, produce a highly entropic distribution of 'FC states' (as shown in the toy example) which means that each individual can have a high frequency of transitions between two or more FC states. Thus, projecting the original data into alternative bases, for instance by applying PCA on the BOLD fMRI time-course, must be handled with care when performing FC state analysis.

4.2. FC state occupancy is a key feature in dFC analysis

Consistent with the findings of [28], we observed that the stationary simulated BOLD time-series from control case I and case III resulted in similar FC states as the ones obtained from the experimental BOLD signal. This similarity may be due to the fact that in these negative controls, the average covariance of each subject is used to synthesize the stationary data. However, when we excluded PCA from the analysis stream, the stationary nature of the simulated data was immediately revealed, since each (simulated) subject tended to classify exclusively into one FC state. That is, all sliding covariance matrices from a simulated subject tend to be classified into the FC state closest to the average covariance of the (actual) subject in question. In contrast, the FC state occupancy for each individual subject is entropic for the experimental BOLD signal and positive control case II, with nonstationary simulated BOLD signals.

4.3. Dynamic reorganization of RSN interactions during resting wakefulness

With all methodological considerations taken into account, our analyses suggest the transition among at least two FC states in resting wakefulness marked by differences in DAN-VIS interactions. Specifically, connections between DAN and VIS networks in individuals during resting-state transition between modes of high and low dFC. These fluctuations are compatible with several prior characterizations of FC, including eye movement associated with waxing and waning outward attention [55, 56], or the modulation of VIS by DAN [57, 58] potentially due to subjects transitioning between wakefulness and light sleep. While further exploration is warranted to fully characterize these relationships, they are indicative of the potential prevalence of dynamic fluctuations in functional connectivity, thus highlighting the importance of such analytical methods in future studies.

5. Appendix

5.1. Toy example

In the toy example of Figure 5, the input white noise first passed through an infinite impulse response (IIR) filter with z-transform in the form of $\sum_{i=0}^p b(i)z^{-i} / \sum_{i=0}^q a(i)z^{-i}$. The coefficients of the IIR filter for f_1 are as follows:

$$\begin{aligned} \mathbf{a}_1 &= [1.0000 \ 0.3907 \ 1.0778 \ -0.2551 \ 0.2416 \ -0.3022] \\ \mathbf{b}_1 &= [0.0673 \ 0.3364 \ 0.6728 \ 0.6728 \ 0.3364 \ 0.0673]. \end{aligned}$$

This filter acts as a lowpass filter with cutoff frequency of 0.15Hz. The coefficients of the IIR filter for f_2 are as follows:

$$\begin{aligned} \mathbf{a}_2 &= [1.0000 \ -1.7391 \ 3.4311 \ -4.0680 \ 5.4505 \ -4.8584 \\ &\quad 4.7419 \ -3.0825 \ 2.2582 \ -0.9128 \ 0.4543] \\ \mathbf{b}_2 &= [0.0082 \ 0 \ -0.0409 \ 0 \ 0.0817 \ 0 \ -0.0817 \ 0 \ 0.0409 \ 0 \ -0.0082]. \end{aligned}$$

This filter acts as a bandpass filter in the range of [0.05, 0.15]Hz. The filtered data are then constrained to have a prespecified covariance matrix. Different covariance matrices were used for systems f_1 and f_2 which are as follows:

$$\text{Cov}_{f_1} = \begin{bmatrix} 1.54 & 2.40 \\ 2.40 & 5.67 \end{bmatrix}, \text{Cov}_{f_2} = \begin{bmatrix} 2.61 & 2.02 \\ 2.02 & 5.36 \end{bmatrix}.$$

5.2. Statistical analysis of occupancy

The results of the statistical test on occupancy of a particular FC state (without loss of generality FC state 1) with and without applying PCA on the original time-series data for different cases are as follows: (K-S test for the experimental BOLD case, $P=0.111$, $\text{stat}=0.5$), (K-S test for the case I, $P=0.031$, $\text{stat}=0.6$), (K-S test for the case II, $P=0.007$, $\text{stat}=0.7$), (K-S test for the case III, $P=0.031$, $\text{stat}=0.6$).

5.3. Static functional connectivity of each subjects

Figure 6 demonstrates the static FC of each subject, calculated over the entire scan time.

5.4. Non-stationary simulation of the BOLD time-series

To simulate a non-stationary BOLD time-series data, we use the results of FC state analysis on the experimental BOLD signal. After applying FC state analysis to the experimental BOLD signal, we fixed the FC state index for each sliding window for all subjects. Then for all sliding windows which are assigned to the first FC state, we calculate the average covariance matrix for all subjects individually. This analysis results in Figure 7A which shows the first FC states for different subjects. In a similar way, we can find covariance matrix associated with the second FC state for each subject (Figure 7B). We then use these covariance matrices to simulate the BOLD signal with similar FC state transitions as the experimental BOLD signal for each subject.

5.5. Mean power spectrum and mean correlation matrix across all the subjects

Figure 8 illustrates the mean correlation matrix and mean power spectrum of all subjects in our dataset.

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Highlights

- Limitations and potential confounds of dimensionality reduction in extracting dFC states from fMRI recordings are evaluated.
- It is shown that the decorrelating actions of principal component analysis can lead to spurious FC state dynamics.
- This limitation is characterized by analyzing the composition and occupancy of FC states in positive and negative controls.
- Accounting for this limitation, two interpretable FC states are extracted during resting wakefulness.

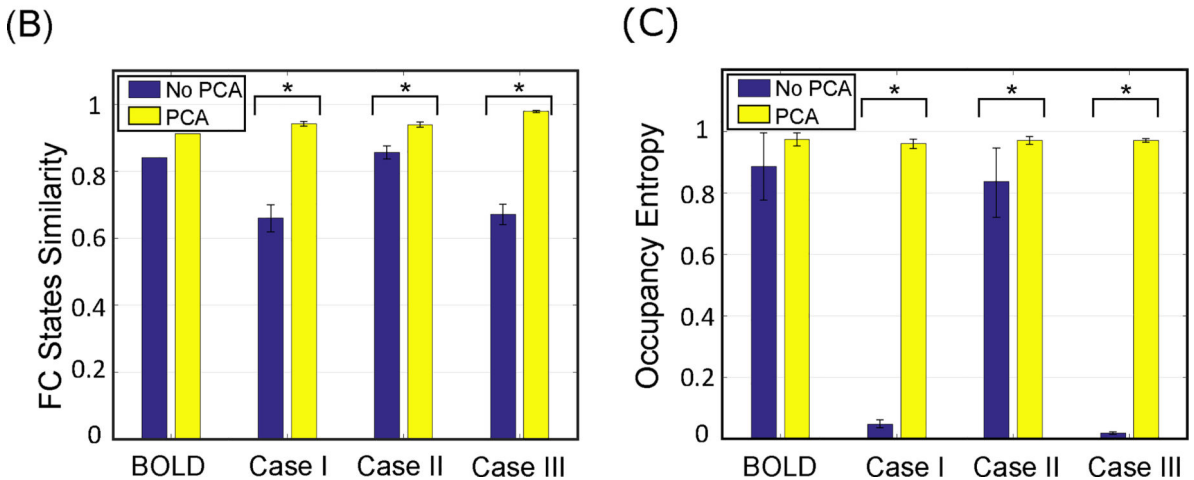
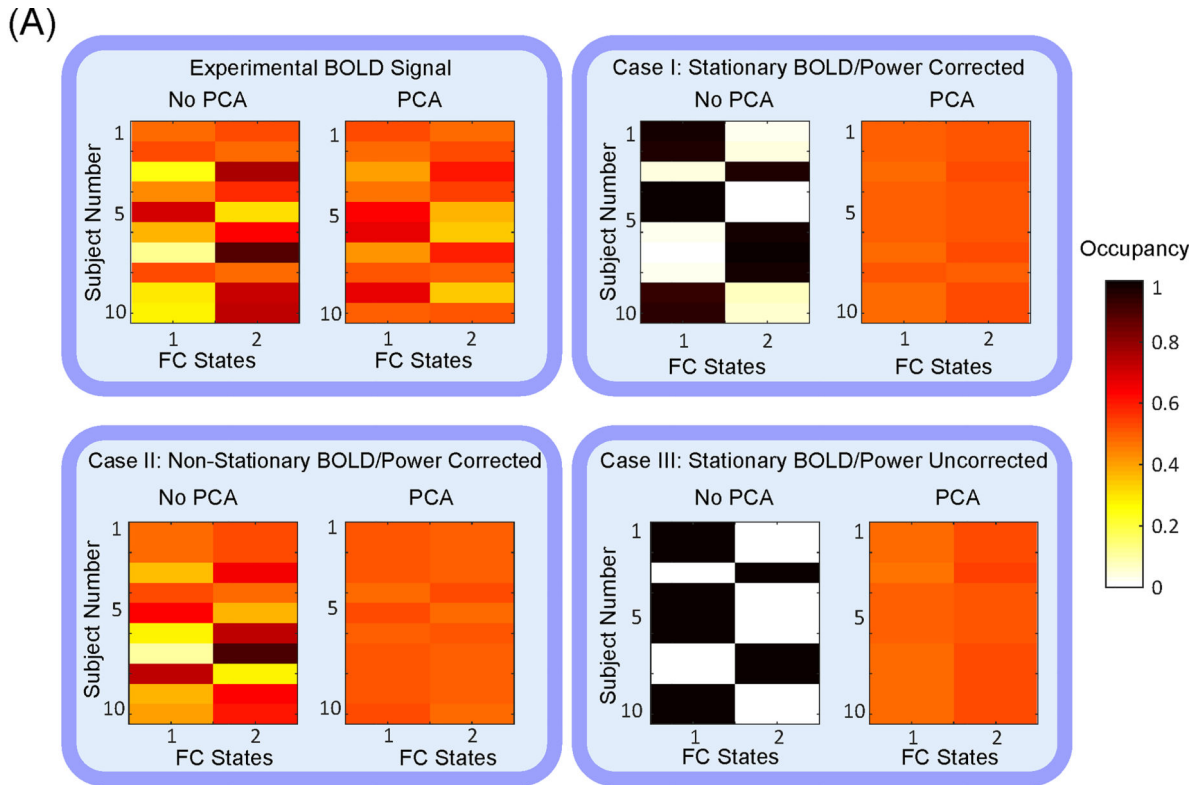
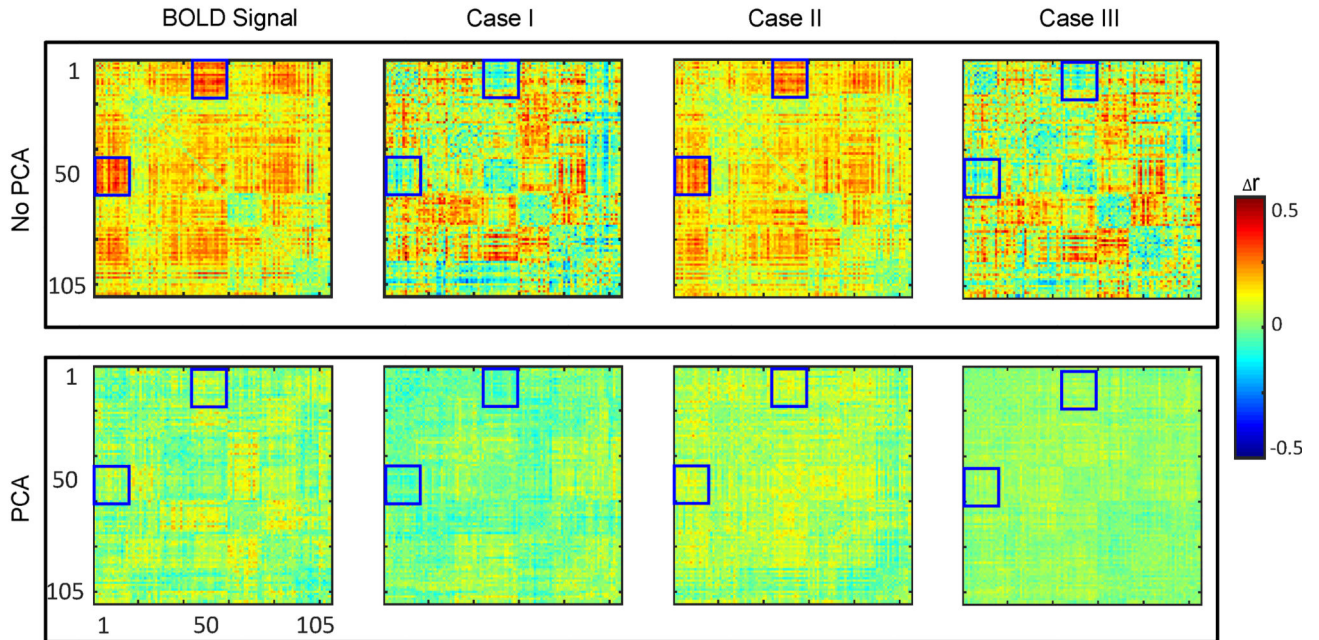


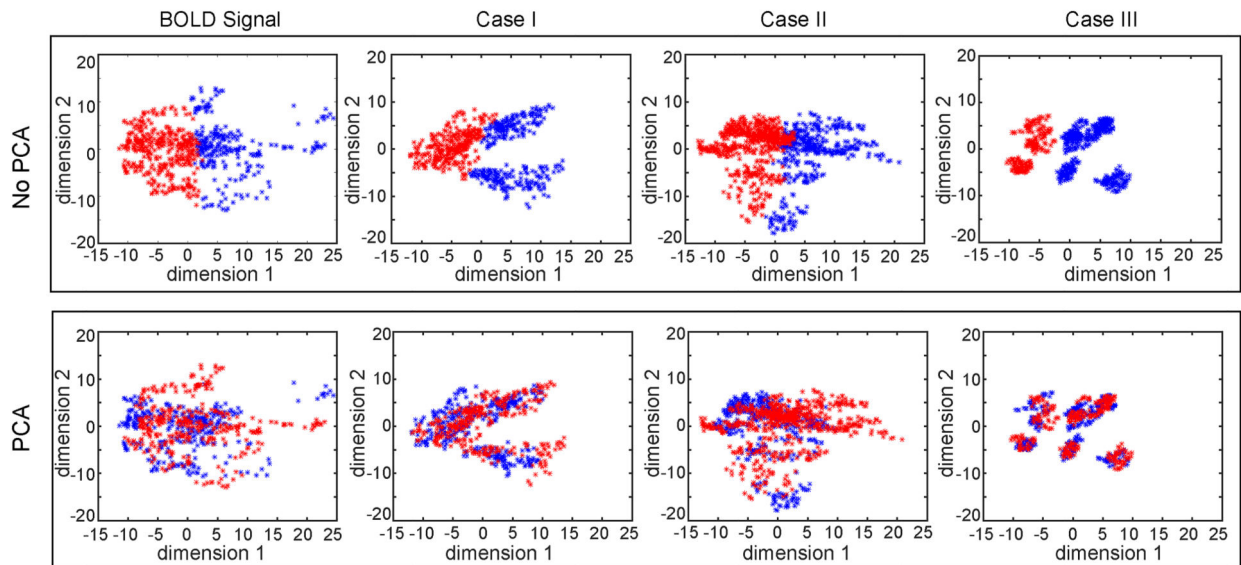
Figure 2.

(A) FC state occupancy across different subject for the experimental BOLD signal, cases I (stationary simulated BOLD, power spectrum corrected), case II (non-stationary simulated BOLD, power spectrum corrected), and case III (stationary simulated BOLD, power spectrum uncorrected) with and without applying PCA to original time-series data. (B) FC state similarity (correlation between two FC states) and (C) FC state entropy for the experimental BOLD, cases I-III with and without applying PCA to original time-series data. Error bars stand for standard deviation (SD).

(A) Difference in FC states (FCS1-FCS2)



(B) 2D visualization of sliding window correlation

**Figure 3.**

(A) Difference between two FC states, separately, obtained from the experimental BOLD signal and simulated BOLD in cases I (stationary simulated BOLD, power spectrum corrected), case II (non-stationary simulated BOLD, power spectrum corrected), and case III (stationary simulated BOLD, power spectrum uncorrected). Top panel shows FC states difference without PCA, while bottom panel represents FC states difference with PCA application. Blue squares represent interaction between VIS and DAN networks. (B) Sliding window correlations projected onto the first two principal components colored by FC state identity. Top panel shows sliding window correlations without PCA, while bottom panel

represents FC states sliding window correlations with PCA application to original time-series data.

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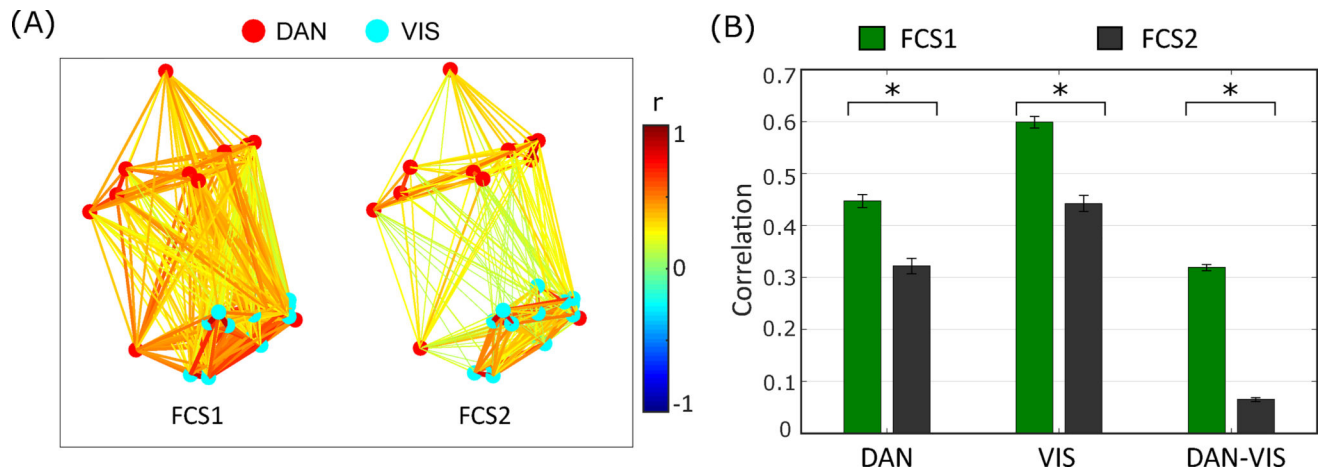


Figure 4.

Connectivity patterns of the two FC states for DAN and VIS networks. All connections with correlation value less than 0.1 are set to zero. (B) Mean correlation connectivity within DAN, within VIS, and between DAN and VIS, for FC states 1 and 2. Error bars represent standard error of the mean.

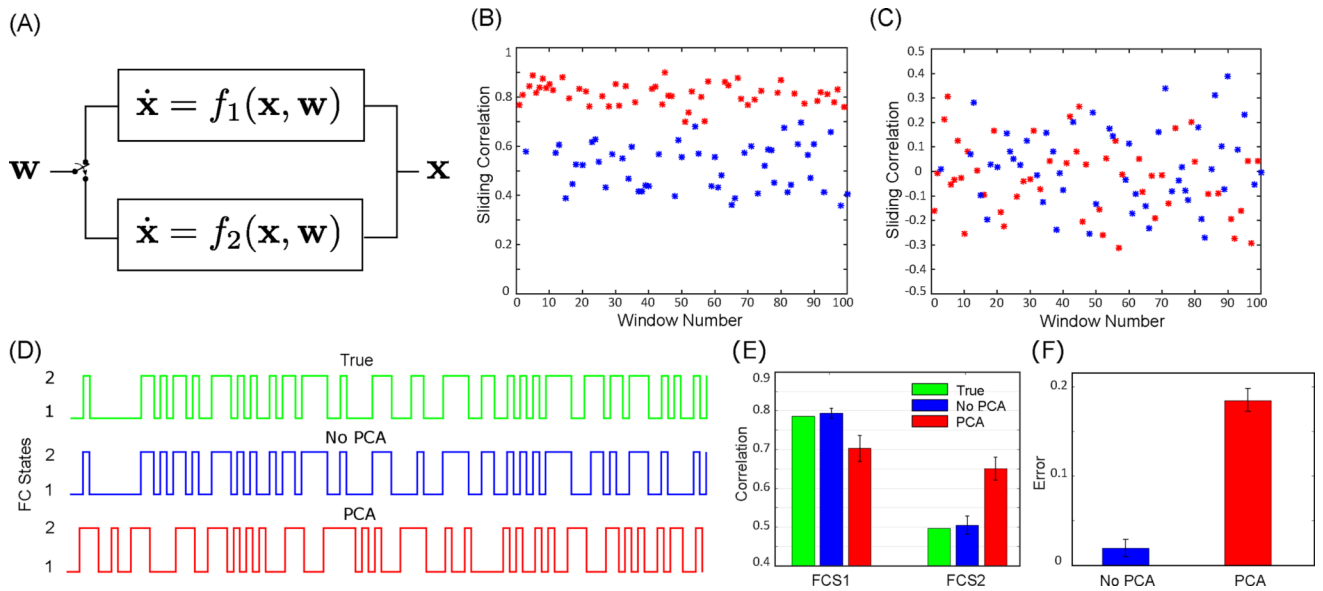


Figure 5.

(A) A two dimensional non-stationary data, \mathbf{x} , randomly generated using two different dynamical systems f_1 and f_2 (red and blue colors) over time. A switch operating randomly determines which dynamical system processes the input white noise, \mathbf{w} , at each time instant. Sliding correlation analysis of the synthesized data (B) before and (C) after applying PCA. (D) (Green) True FC state of the synthesized data chosen by the switch operating randomly. (Blue) Estimated FC states with application of PCA to the synthesized data. (Red) Estimated FC states with application of PCA to the synthesized data. (E) Results of FC state analysis for estimating correlation between two time-series in both FC states with and without PCA over 100 realization of random synthesized data. (F) Estimated correlation error with and without applying PCA to time-series data for both FC states. Error bars stand for standard deviation (SD).

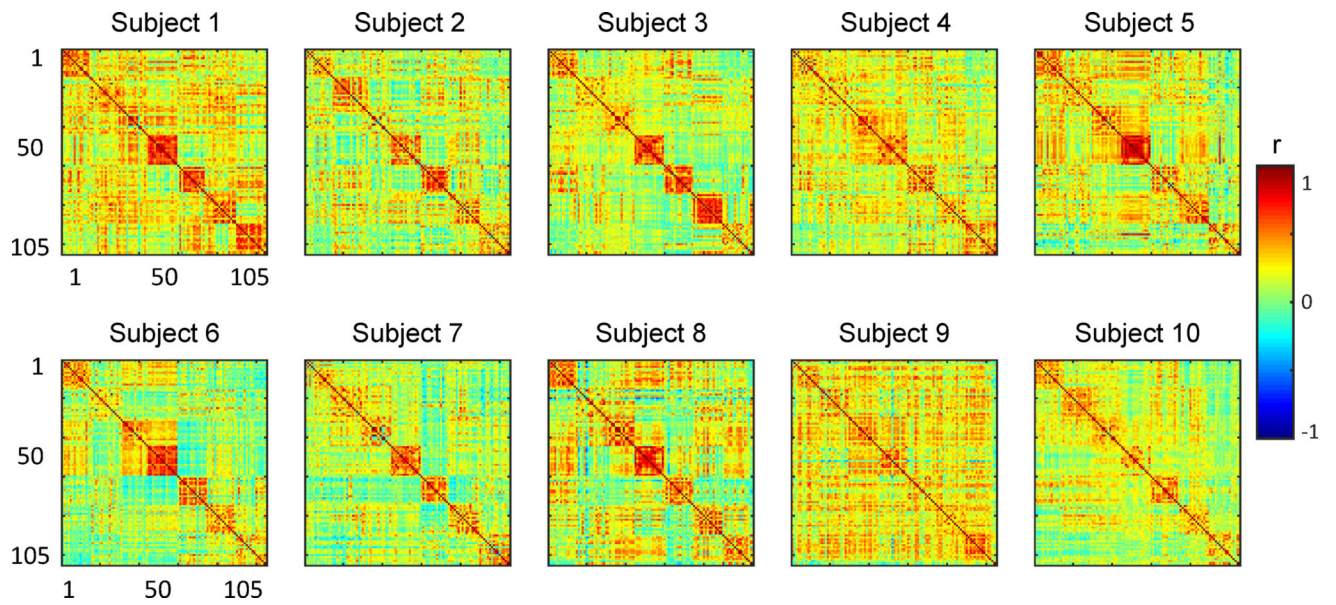


Figure 6. Static FC of each subjects, calculated over the entire scan time, for all individuals in our dataset.

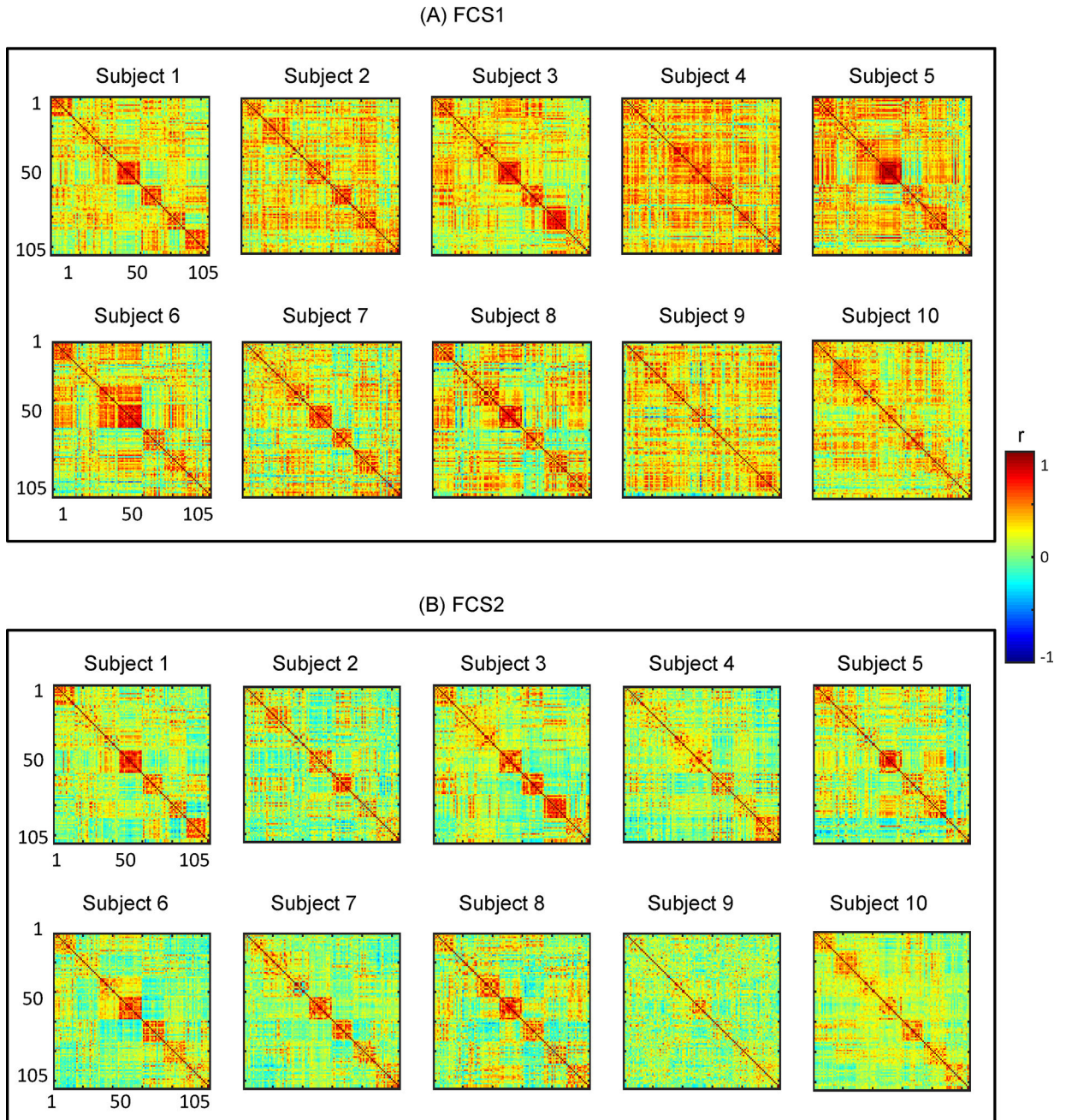


Figure 7.

FC states used for simulation of time-series data in case II. (A) Correlation matrices associated to the first FC state for each individual obtained from FC state analysis of the BOLD signal. (B) Correlation matrices associated to the second FC state for each individual obtained from FC state analysis of the BOLD signal.

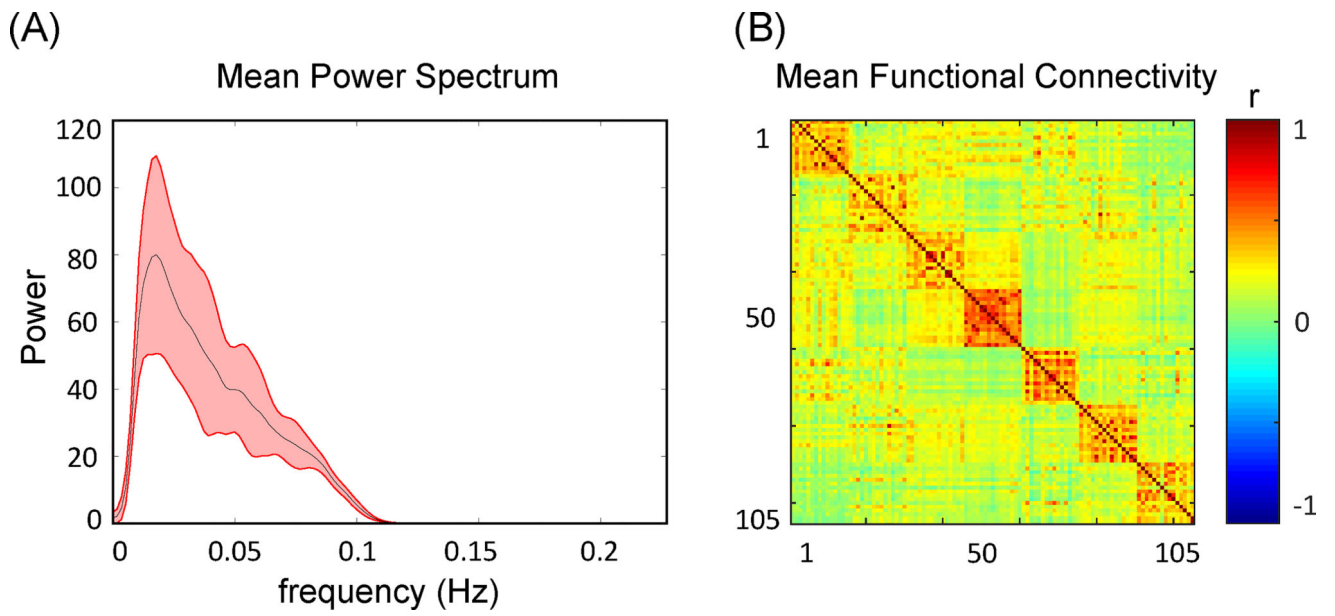


Figure 8. (A) Mean power spectrum and (B) mean static functional connectivity of all subjects in our dataset.