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Distributed patterns of brain activity underlying real-time fMRI neurofeedback training

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Abstract—Neurofeedback (NF) based on real-time functional magnetic resonance imaging (rt-fMRI) is an exciting neuroimaging application. In most rt-fMRI NF studies, the activity level of a single region of interest (ROI) is provided as a feedback signal, which the participants are trained to up or down-regulate. Neurofeedback training effects are typically analyzed using a confirmatory univariate approach, i.e., changes in the target ROI are explained by a univariate linear modulation. However, learning to self-regulate ROI activity through NF is mediated by distributed changes across the brain. Here we deploy a multivariate decoding model for assessing NF training effects across the whole brain. Specifically, we first explain the NF training effect by a post-hoc multivariate model that leads to a pattern of co-activation based on 90 functional atlas regions. We then use cross-validation to reveal the set of brain regions with the best fit. This novel approach was applied to data from a rt-fMRI NF study where participants learned to down-regulate the auditory cortex. We found that the optimal model consisted of 16 brain regions whose co-activation patterns best described the training effect over the NF training days. Cross-validation of the multivariate model showed that it generalized across participants. Interestingly, the participants could be clustered into two groups with distinct patterns of co-activation, potentially reflecting different NF learning strategies. Overall, our findings revealed that multiple brain regions are involved in learning to regulate activity in a single ROI, and thus lead to a better understanding of the mechanisms underlying NF training.

Index Terms—real-time fMRI, neurofeedback, functional networks, co-activation, basal-ganglia, thalamus, insula

I. INTRODUCTION

Neurofeedback (NF) based on real-time functional magnetic resonance imaging (rt-fMRI) is an emerging technique that allows to train participants voluntary control over their own brain activity [26], [25]. In the majority of rt-fMRI NF experiments, the feedback signal reflects neuronal activity within a single region-of-interest (ROI), which participants are taught to up or down-regulate. Previous rt-fMRI NF training studies have demonstrated that healthy participants can indeed gain control over localized brain activity, and that such training affects behavior. For example, training of the parahippocampal cortex modulated memory function [77], of the right anterior

cingulate cortex (ACC) reduced pain perception [62], [22], of the precentral gyrus speeded up motor responses [31], of the inferior frontal gyrus improved linguistic performance [1], of the insula modulated emotions [5], of the occipital cortex improved visual perception [29], [33], and of the right auditory cortex modulated auditory perception [19], [9]. Recent studies have also demonstrated clinical relevance of rt-fMRI-based NF training. For example, chronic pain patients were trained to regulate the ACC [62], chronic tinnitus patients learned control over the auditory cortex [20], Parkinson's disease patients learned control over the supplementary motor area [10], major depression patients learned to increase activity in brain regions involved in positive emotions [28], chronic stroke patients learned control over the ventral premotor cortex [28], nicotine addicts learned control over the ACC and ROIs in the prefrontal cortex [34], and schizophrenia patients learned control over the insular cortex [15].

To shed light on the neural underpinnings of successful self-regulation, training-related changes in brain networks have been investigated post-hoc for some of the above-mentioned ROI-based NF studies. For example, Rota et al. analyzed functional connectivity (FC) by using a seed correlation approach that revealed FC changes with the NF target ROI (i.e., the inferior frontal gyrus) as a function of NF training [16], [1]. Their FC analyses revealed changes in pairwise correlations between the NF target ROI and other brain regions, but FC changes between brain regions other than the NF target ROI could not be detected with the seed correlation approach. Ruiz et al., studied changes in effective connectivity that were associated with NF training of the insular cortex using Granger causality modeling (GCM), which is a connectivity analysis approach in which time-series from pre-selected brain regions are used to predict time courses of another region [15] [3], [12], [13]. They found enhanced effective connectivity of the NF target ROI with other brain areas, but similar as with the seed-based approach, Granger causality limits the number of brain regions whose connectivity changes were analyzed. Scharnowski et al. found post-hoc connectivity changes associated with NF training of the visual cortex using

a psychophysiological interaction (PPI) analysis, and dynamic causal modeling (DCM) [33], [69]. Whereas PPI is exploratory and allows to identify task-related correlation changes with the NF target ROI [70], DCM is hypothesis-driven and uses Bayesian model comparison to compare which network architecture explains the data best [72]. However, similar to the seed-based approach, PPI allows to analyze only FC changes with the NF target ROI, and similar to Granger causality, DCM allows to analyze effective connectivity of only a few pre-defined ROIs. Finally, Haller et al. applied independent component analysis (ICA) to reveal network changes associated with training the auditory cortex [19], [9]. Their analysis revealed functionally relevant independent components (ICs), including the auditory network that contained the NF target ROI, the default mode network (DMN), and the executive control network. Even though this study used a data-driven multivariate approach for the decomposition into ICs, the FC analysis was done using a univariate approach (i.e., changes in pairwise correlations between IC timecourses over training days were analyzed separately). Finally, a recent study by Harmelech et al. demonstrated that even a single session of ROI-based NF training induced lasting changes of FC within the DMN [27]. Overall, these post-hoc analyses of global changes associated with ROI-based NF training indicate that the effects of NF training extend beyond the target ROI. However, these investigations focused on either hypothesis-driven multivariate analyses observing connectivity changes between the target ROI and a limited number of regions, or of whole brain univariate analyses observing changes in pairwise connections. Here we extend these previous approaches by proposing a post-hoc multivariate decoding method to identify changes in brain activity across the whole brain that are associated with NF training. This decoding analysis includes activity signals from all brain regions, which are used to explain NF training effects in a multivariate model. Specifically, we (a) used a linear model, but with a multivariate decoding setup where time courses of all brain regions as regressors, and the interaction between the self-regulation paradigm and a linear improvement across NF training as the target signal; (b) ranked all brain regions according to their consistency over subjects; (c) test this hypothesis include also an estimation of the optimal model size by implementing a leave-one-participant-out (LOO) cross-validation and backward elimination methods.

Our new multivariate approach will decode and identify co-activation maps of brain regions involved in improved self-regulation skills, and will thus reveal the co-activation patterns underlying ROI-based NF training. To validate the method, we applied this novel approach to data from a previous ROI-based NF study, in which 12 participants learned down-regulation of the right auditory cortex activity while being presented with acoustic stimulation [19], [9].

II. MATERIALS AND METHODS

A. Data description

Details about the participants, task procedure, and data acquisition can be found in [19], [9]. For completeness, the main parameters are summarized here.

Participants

Twelve healthy right-handed volunteers with normal audition took part in the study. Mean age of participants was 28.37 years (range 24-33 years). The study was approved by the local ethics committee, and all participants gave written informed consent. Before the experiment, volunteers received written instructions explaining that they will learn to down-regulate their primary auditory cortex activity with the help of NF.

Task procedure

Each participant had four days (sessions) of NF training with approximately 1 week intervals between them. At each day, participants had four runs, which led to a total of 16 runs per participant. Before each training session, the participants primary auditory area was identified using a standard fMRI auditory block-design paradigm, consisting of 20 s ON and 20 s OFF bilateral auditory stimulation using a 1000 Hz pulsating sine tone, repeated five times. Following the localizer, four rt-fMRI NF training runs were performed, the right localized auditory region was the target ROI. Each run was composed of five 30 s baseline blocks, interleaved with 60 s down-regulation blocks. The same pulsating sine tone of 1000 Hz was provided as auditory stimulation during down-regulation blocks. The signal from the target ROI was provided as a visual feedback during the entire run as a moving line graph. The participants were informed about the data processing delay of about 1 s and of the intrinsic physiological hemodynamic response delay of about 6 s during the down-regulation periods. No specific regulation strategy was recommended to the participants, but it was emphasized that they should find an individual strategy that worked best for them. For online data analysis and feedback presentation we used the Turbo Brain-Voyager software package (Brain Innovation, Maastricht, The Netherlands) in combination with in-house Matlab (Mathworks Inc., Natick, MA, USA) scripts.

Data acquisition

The experiment was performed on a 3 T whole-body MR scanner with a standard 12-channel head coil (Siemens Magnetom Verio, Siemens Erlangen, Germany). Functional data were acquired with a Echo-Planar Imaging (EPI) sequence (echo time (TE) 40 ms, repetition time (TR) 2000 ms, matrix size 64×64 , voxel size $3 \times 3 \times 3 \text{ mm}^3$, and 19 repetitions). Additionally, we acquired an anatomical T1-weighted whole brain image using a Magnetization Prepared Rapid Gradient Echo (MPRAGE) sequence (matrix size 256×256 , 176 partitions, 1 mm^3 isotropic voxels, 26 slices with 1 mm thickness).

B. Data preprocessing

Preprocessing was performed using SPM8 software (Wellcome Department of Imaging Neuroscience, London, UK). All functional volumes were spatially realigned to the first volume of each run and normalized into MNI space (Montreal Neurological Institute, resampled voxel size: $2 \times 2 \times 2 \text{ mm}$) by using cubic B-spline interpolation. Image series were then parcellated into $V = 90$ regions based on the Greicius functional atlas [11], and time courses were regionally averaged, de-meaned and linearly de-trended using Matlab standard

functions. For each participant n , data was concatenated into a single matrix for all 16 runs: $\bar{Y}^{(n)} = [Y_1^{(n)} Y_2^{(n)} \dots Y_{16}^{(n)}]$, where $Y_k^{(n)}$ is the spatio-temporal matrix of run k .

C. Within-subject model

We deploy a linear model to reveal main effect of regulation. For each participant n time courses $\bar{Y}^{(n)}$ are used to explain the block paradigm Z_{block} . Z_{block} is a vector of length $16 \cdot 195$, constructed by 16 concatenated block designs, normalized to zero mean and unit variance (Fig. 1). This linear model is defined for each participant n as follows:

$$Z_{block} = \bar{Y}^{(n)T} \cdot \beta_{main}^{(n)} + \epsilon_{main}^{(n)}, \quad (1)$$

where $\beta_{main}^{(n)}$ is the parameter vector and $\epsilon_{main}^{(n)}$ is the noise term. The optimal parameter estimates $\hat{\beta}_{main}^{(n)}$ are found by minimizing the sums of squares of the residuals $\hat{\epsilon}_{main}^{(n)} = Z_{block} - \bar{Y}^{(n)T} \times \hat{\beta}_{main}^{(n)}$ between the predicted and the fitted model, which is optimal assuming $\epsilon_{main}^{(n)}$ is i.i.d. normally distributed.

Next, we deployed another linear model to reveal the training effect over days, here the time courses $\bar{Y}^{(n)}$ are used explain the interaction between the linear improvement across training days and the block-design paradigm. For this purpose we defined the improvement signal:

$$Z_{target} = Z_{training} \odot Z_{block} \quad (2)$$

where $Z_{training}^T = \underbrace{[3 \dots 3]}_{4 \cdot 195} \underbrace{[1 \dots 1]}_{4 \cdot 195} \underbrace{[-1 \dots -1]}_{4 \cdot 195} \underbrace{[-3 \dots -3]}_{4 \cdot 195}$. $Z_{training}$ is also normalized to unit variance, and then multiplied element-wise (\odot) with Z_{block} to generate the improvement signal Z_{target} (Fig. 1). The linear model for each participant n was defined as follows:

$$Z_{target} = \bar{Y}^{(n)T} \cdot \beta^{(n)} + \epsilon^{(n)}, \quad (3)$$

where $\beta^{(n)}$ is the parameter vector and $\epsilon^{(n)}$ is the noise term.

To test the null hypothesis that the improvement signal is explained better than by chance, we phase-randomized the matrix $\bar{Y}^{(n)}$ (along its rows) in the temporal Fourier domain. The null is rejected if the estimation error for the real data is significantly better than for surrogate data. Specifically, generating 19 surrogate data sets allows significance to be established at 5%. Participants that fit the model well are those with real-data residual sum of squares (RSS) values lower than all 19 surrogates RSS values (confidence intervals of 95%). For future use, we defined a $195 \times |\mathcal{N}|$ matrix based on a general set \mathcal{N} : $\hat{\beta}_{\mathcal{N}}^+ = [\dots \hat{\beta}^j \dots]$, $j \in \mathcal{N}$. According to the RSS single-participant performances, we defined \mathcal{M} as the set of “good” participants that successfully fitted the model, and the estimated parameter matrix $\hat{\beta}_{\mathcal{M}}^+$ accordingly.

D. Regions involved in improvement of self regulation

In order to investigate which are the key regions for improving of self-regulation; we analyzed $\hat{\beta}_{\mathcal{M}}^+$, by performing a two-sided one-sample t-test on each row (i.e., per brain

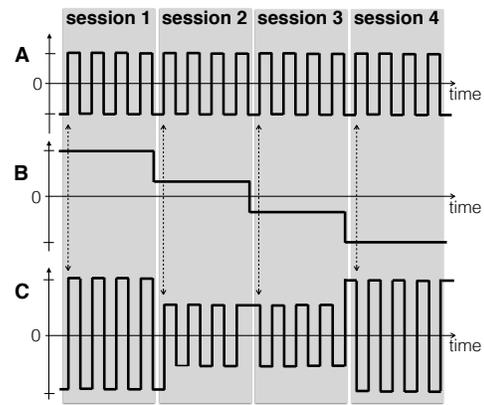


Fig. 1. Construction of the Z_{target} signal. (A) Z_{block} block paradigm of all runs concatenated in time, indicating the regulation and rest periods. (B) $Z_{training}$ a linear descending training signal over four training days. (C) Z_{target} element-wise multiplication between $Z_{training}$ and Z_{block} .

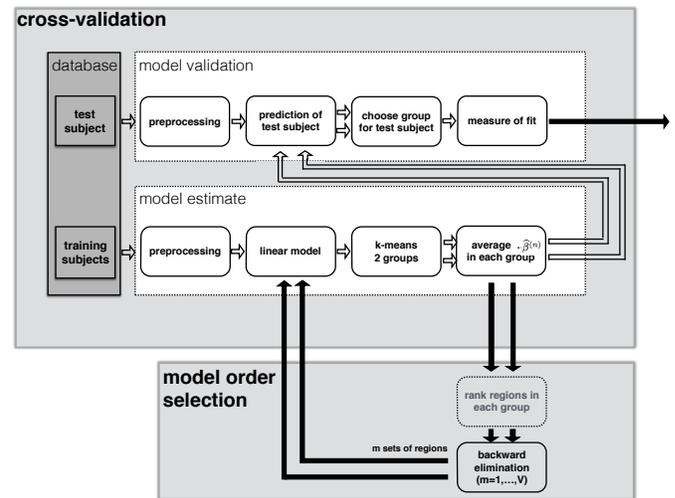


Fig. 2. Overview of the processing pipe-line with cross-validation. The model estimation was done for all participants but one (test participant). The model validation for test participant used the model estimation results. Backward elimination for brain regions repeats this procedure for all possible model sizes. Regions were eliminated based on the ranking of each set which was calculated once using the full model (i.e., $m = V$). Results are the measure of fit calculated for each possible model order size.

region across participants). Based on the results, we ranked the regions according to their t-values. We also included the anatomical location of the functional network regions (Table I). Next, the generalizability of the model across participants was tested with a leave-one-participant-out (LOO) cross-validation scheme (Fig. 2). In this approach, the fitted model for $|\mathcal{M}| - 1$ training participants was used to establish the corresponding $\hat{\beta}$, that was then applied to the left-out participant. In each LOO iteration, we selected a left-out participant n , defined the training set $\mathcal{M}^{(n)} = \mathcal{M} \setminus \{n\}$, and updated matrix $\hat{\beta}_{\mathcal{M}^{(n)}}^+$ accordingly. Due to the high inter-participant variability, we clustered $\hat{\beta}_{\mathcal{M}^{(n)}}^+$ into two groups using k-means clustering that used the cosine distance measure with 10'000 replicates, resulting in two sets of participants: $\mathcal{M}_i^{(n)} = \{j : j \in \mathcal{M}^{(n)} | j \in \text{group } i\}$, $i = 1, 2$. To avoid double dipping, k-mean clustering was calculated within each fold and the number of clusters was consistently set to two

based on the Calinski-Harabasz criterion [71]. For each group i , we adjusted the fitted data matrix $\hat{\beta}_{\mathcal{M}_i^{(n)}}^+$, based on which we calculated the training response:

$$\bar{\beta}_{\mathcal{M}_i^{(n)}} = 1/(|\mathcal{M}_i^{(n)}|) \cdot \hat{\beta}_{\mathcal{M}_i^{(n)}}^+ \cdot \mathbf{1}, \quad (4)$$

where $\mathbf{1}$ is a one vector of length $|\mathcal{M}_i^{(n)}|$. The prediction for the left-out participant n was defined as:

$$\bar{Z}_i^{(n)} = \bar{Y}^{(n)T} \cdot \bar{\beta}_{\mathcal{M}_i^{(n)}} \quad (5)$$

Furthermore, we defined the measure of fit as the correlation between improvement paradigm Z_{target} and the prediction signal

$$\bar{F}_i^{(n)} = \text{corr}(Z_{target}, \bar{Z}_i^{(n)}) \quad (6)$$

The group level co-activation maps were calculated separately for each of the two groups using the following steps:

- (1) rankings $\bar{\beta}_{\mathcal{M}_i^{(n)}}$ according to one-sample t-test were calculated for each group in each fold;
- (2) the optimal model (top 16 regions, see next paragraph) in each fold was identified;
- (3) a list counting the times each region was part of the optimal model was created;
- (4) co-activations maps were defined based on the list created in step (3), regions that were ranked within the top 16 regions for more than 6 out of 11 folds were selected.

E. Model order selection

Region ranking for optimal model size analysis was calculated once for each LOO fold. A one-sample t-test was calculated for the training set matrix $\bar{\beta}_{\mathcal{M}_i^{(n)}}$ that included all regions (i.e., full model $m = V$), results were sorted in descending order which set the ranking of regions. The search for the optimal model order was done using backwards elimination approach [43] that was executed separately for each fold, where at the initial step all regions are included and in following steps regions are eliminated based on the ranking that was determined. The performances analysis of all possible model sizes was done separately for each fold by exploring the model estimation of the training set (i.e., two group clustering and $\bar{\beta}_{\mathcal{M}_i^{(n)}}$ in each of the groups) and model validation (i.e., the measure of fit $\bar{F}_{i,m}^{(n)}$, Eq. 6). Based on those results we could assign the left-out participant n to one of the two clusters $\mathcal{M}_i^{(n)}$, $i = 1, 2$ by maximizing the mean measure of fit between the two clusters: $\{n \in \mathcal{M}_k^{(n)} \mid \sum_{j=1}^V \bar{F}_{k,j}^{(n)} > \sum_{j=1}^V \bar{F}_{t,j}^{(n)}\}$. The performance results of all the folds were consolidated by averaging the measure of fit over all folds, resulting in one measure for each possible model size. The selection of the optimal model size was based on comparing that average measure of fit to the performances of 1000 surrogate data sets, and choosing the order size that maximizes the data performance and exceed the 98th percentile of the surrogate distribution.

III. RESULTS

A. Learned down-regulation of primary auditory cortex

As reported previously, over the course of four days of NF training, participants learned to down-regulate activity in the NF target region, i.e. the right auditory cortex [19], [9].

During the experiment, feedback was only based on the activity level of a single subject-specific target single ROI determined using a functional localizer. Here we investigated training effects based on combined activity from multiple brain regions that were taken from a predefined functional atlas.

B. Within-subject discriminative model

In the analysis of the main effect we found positive bilateral activation in the auditory regions and frontal medial cortex, contralateral thalamic activity, caudate and middle occipital lobe (left regions with activations while right regions with deactivation); and bilateral deactivation of insula, medial superior frontal gyrus and calcarine. Next, for the training effect analysis we found that all except one participant showed significant training effect as function of training days (Fig.

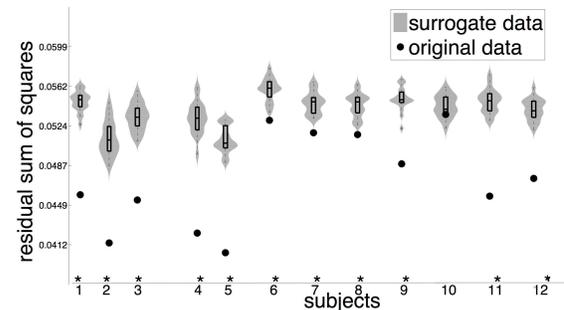


Fig. 3. RSS for all participants. RSS of the original data (circles) was significantly lower than the RSS of the surrogate data (\diamond) for eight participants (1, 2, 3, 4, 5, 9, 11, 12; indicated by *). Participants 6,7,8 present low performances but still within the 95% significant levels (also indicated by *). One participant (number 10) did not show a significant learning effect across NF training.

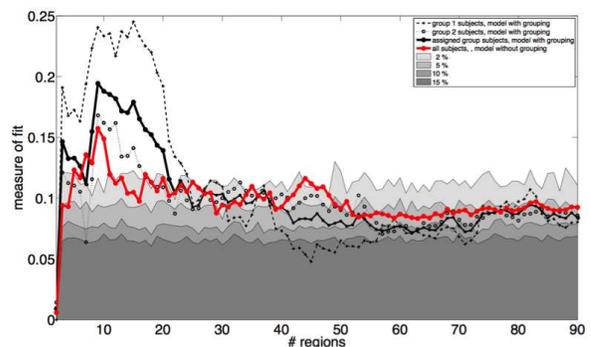


Fig. 5. Performance of cross-validation over different model sizes. Measure of fit indicate average group level correlation between Z_{target} and test participants prediction. Dashed line represents the results for group 1, dotted line for group 2; solid line for assigned group; i.e., each test participant was assigned to the group with the best fit; red line without subdividing into two groups; i.e., for each left-out subject, the model is estimated based on all other subjects. Gray scale represent the percentile of the surrogate distribution.

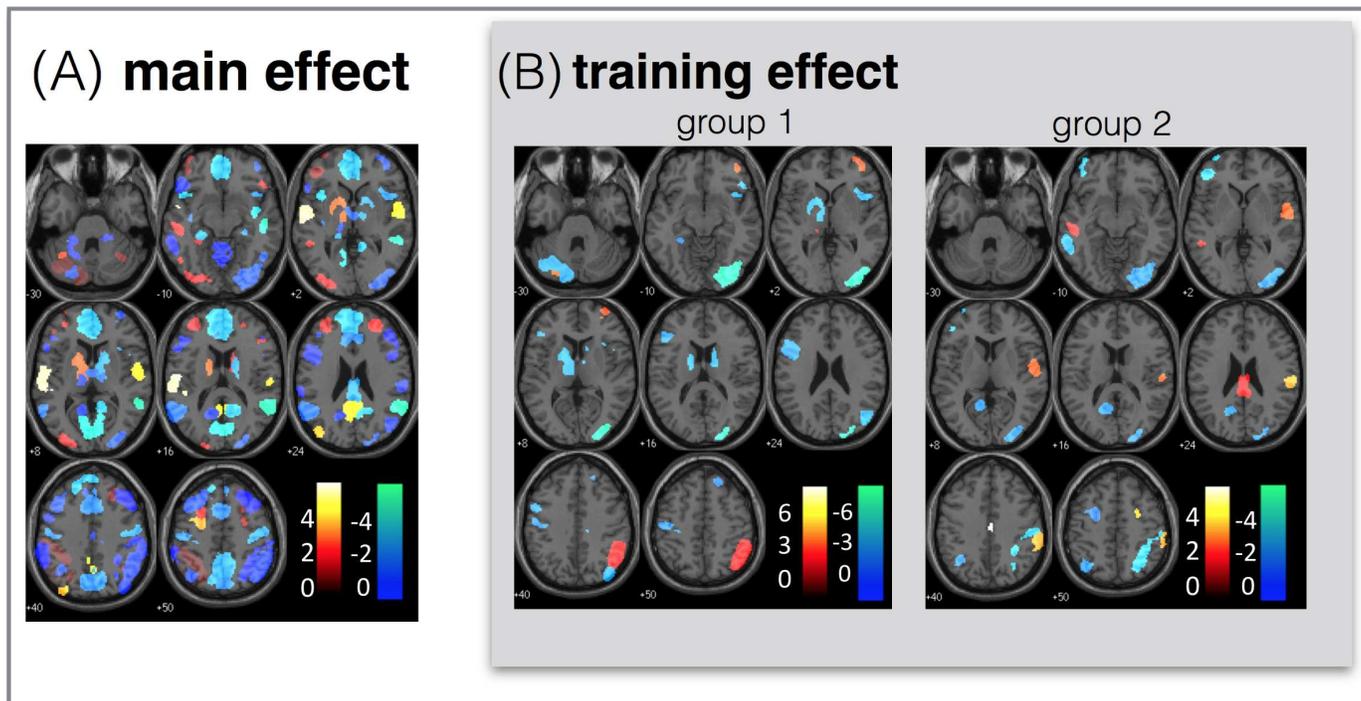


Fig. 4. Co-activation maps shown as axial slices in neurological convention. (A) T-value of the main effect analysis for all brain regions. (B) Optimal model for the rt-fMRI training effect over 4 sessions. Shown here are the positive and negative t-values averaged across the within-subject multivariate models.

3) $\mathcal{M} = \{1, 2 \dots 9, 11, 12\}$. For those who did, the linear combination of activity in the V regions was related to changes in regulation strength across NF days. Table I shows the brain regions ranked according to the t-values and also reports the average linear combination values (i.e. β values).

C. Regions involved in improving self-regulation

A complete view of the learning effect needs to consider two results, the first is the main effect that corresponds to the average activation level during down regulation session and the second is the change of activation over training sessions. The multivariate analysis of the training shows the co-activated regions that explain a linear change in activation over sessions. The clustering analysis revealed two distinct groups of participants. Cross-validation identified distinct sets of brain regions that were involved in training (see Fig. 4). Here, we examine closer the two groups: the optimal model for group 1 includes left Crus 1, right insula, caudate, right midcingulate cortex, right angular gyrus, right thalamus and Putamen areas that are associated with negative beta values (an increase of activation over the session) while left frontal operculum, left crus I and right middle frontal gyrus are associated with positive beta values (a decrease of activation over the session). In the optimal model for group 2 it was found that right inferior parietal lobule, right angular gyrus, left inferior temporal gyrus and left inferior frontal gyrus are associated with negative beta values and increase of activation, while midcingulate cortex, left middle temporal gyrus, right superior temporal gyrus (i.e., the target auditory ROI) and right supramarginal gyrus are associated with positive beta values and thus decrease of activation.

D. Model order selection

The performance of the proposed model was evaluated using the average measure of fit; i.e., correlation between the improvement paradigm and the prediction for the test participant. Each test participant n was assigned to its best fitted group. In Fig. 5, we present these cross-validation results for the real and surrogate data. The distribution of the test statistic under the null hypothesis indicated that 16 regions can be considered as an optimal model order. **An examination of the optimal models across the folds revealed that the t-values used for the ranking of the top 16 regions were significant ($t - value > 2, p < 0.5$)** It is important to note, that for the optimal model size, the cross-validation results indicated that the two groups analysis with real training data outperformed the analysis including all subjects with real data (i.e., without subdividing subjects in two groups) as well as with surrogate data (Fig. 5). Despite the high inter-participant variability, this further confirmed that the top ranked regions can be generalized to out-of-fold participants.

IV. DISCUSSION

Voluntary control over brain activity in a single ROI can be learned using rt-fMRI NF. Here, we deployed a multivariate data-driven model to reveal how co-activation of multiple brain regions explain successful NF training. For evaluating the consistency of the activated regions across participants, we used cross-validation to determine the most economical and generalizable model, which consisted of 16 brain regions. Our results show that (I) NF training of a single ROI caused distributed changes across the whole-brain; (II) a multivariate

model of co-activated brain regions that generalizes across participants can be identified; and (III) participants can be clustered into two distinct groups who each co-activated different sets of brain regions.

A. Post-hoc analyses of functional network reorganization

Training brain activity in a single ROI using rt-fMRI NF does not only effect the NF target ROI, but also other regions across the brain. A better characterization of these changes is important for understanding the neural underpinnings of NF training, thus potentially improving its efficacy. As presented in the Introduction, the methods that were previously used to investigate global distributed changes include seed-based correlation, whole-brain pairwise correlation, GCM, DCM, and PPI. Each of these approaches has its advantages and disadvantages. For example, GCM and DCM allow to determine the directionality of connectivity changes, and DCM allows for modeling of effective connectivity at the neuronal level [72]. On one hand, these multivariate approaches are limited to analyzing connectivity changes of only a limited number of brain regions that have to be defined a priori. On the other hand, seed-based correlation, whole-brain pairwise correlation, and PPI can handle more brain regions, but they can consider only pairwise connections and therefore cannot detect interactions between multiple brain regions. Our proposed multivariate approach is complementary in that it allows for investigating changes related to NF training by activity traces across the entire brain. This does not require any prior assumption about how brain regions interaction. Such multivariate interactions that are characterized by co-activation patterns and that are specific to the improvement of self-regulation across NF training days could not be revealed by the previously used methods. Therefore our approach extends previous investigations of changes related to NF-training.

Previous analyses were all carried out on the group-level, even though NF learning success, strategies, and its neural underpinning vary substantially between participants [25]. We examined generalization and consistency of the results using cross-validation, as it was implemented in our approach allows ensuring and quantifying how the co-activation maps generalize across participants. To account for the high inter-participant variability of learning in NF experiments, we applied clustering procedures where participants were clustered according to similarities in their co-activation maps. Our analysis revealed two distinct and consistent groups of participants, which showed differences and commonalities in NF learning across participants.

B. Self-regulation of activity in auditory cortex implicates distributed set of regions

When applying the proposed analysis method to data from a NF training study where participants learned to down-regulate the ROI auditory cortex, we found evidence for changes in a distributed set of brain regions that was associated with NF learning. Specifically, we found that a co-activation model with 16 brain regions best explained the NF training effect. The clustering analysis had revealed two groups, each with

a distinct co-activation pattern. In the first group, the co-activation pattern consisted of brain regions mainly related to self-awareness (e.g., precuneus, insula, angular gyrus) [73], [74], [75], cognitive control (e.g., frontal operculum) [76], and skill learning (e.g., caudate nucleus, putamen) [85]. Especially the latter is interesting, because they have frequently been reported to be involved in NF experiments [58], [59], [22], [77], [86], and it has recently been proposed that NF learning is linked to skill learning [78]. In contrast, in the second group, the co-activation pattern consisted of brain regions related to the auditory/language pathway (e.g., right superior temporal gyrus, which contains the target ROI, as well left inferior frontal gyrus, and supramarginal gyrus) [79], [80], sensory information processing (e.g., inferior parietal lobe) [81], [82], and reward-related learning (e.g., midcingulate gyrus) [54][48].

To support the validity of the two group clustering, we have compared the group performances against the one obtained when not separating the subjects into two groups; i.e., for each left-out subject, the model was estimated based on all other subjects. The results suggest that the models for the subgroups were not only different, but also lead to a more accurate model than the one obtained jointly from all subjects. Although interpreting the differences in co-activation patterns between the two subgroups would require extensive meta-analytic profiling, their functional differences already suggest different learning strategies. For example, participants in group 1 might have adopted an explicit skill learning strategy whereas participants in group 2 were more prone to implicit reinforcement- and reward-based learning [65]. Detailed reports of the cognitive strategies that participants used during the NF experiment would be useful to further elaborate this speculation, but no such reports are available for this study. Those co-activations maps could potentially be used for additional purposes, for example, monitor self-regulation aptness. Recent method proposed to detect on-line arousal level using only fMRI data [87] can be adapted for this purpose, co-activations maps could be projected onto on-line rt-fMRI volumes and generating rt-fMRI self-training index.

C. Limitations

The first limitation is that we used a linear model as a first-order approximation of the NF learning effect. Although the cross-validation results confirm that the linear assumption holds, it might have not been the optimal model.

The second limitation is that in this multivariate analysis we included all brain regions, including the right auditory region based on the Greicius atlas. Since the atlas-based definition of the auditory region differs from that of the auditory NF target ROI (which was based on a functional localizer for each participant), these regions are not identical and thus there is no ‘double-dipping’. Including the atlas auditory ROI in the model allows discovering whether it plays a role (or not) in the learning effects of NF.

Finally, our sample size is low for clustering the participants into two groups. However, our unsupervised clustering showed a clear separation into two groups, and despite the lower

sample size per group, the cross-validation results are superior in the subgroups compared to all subjects combined. Since inter-subject variability in training strategies is an important topic in neurofeedback training studies, future rt-fMRI studies in larger samples might use similar clustering approaches to identify different learning strategies.

D. Conclusion

The proposed multivariate approach revealed interactions between distributed brain regions that contributed to learning control over a ROI through NF training. Using a cross-validation scheme, we examined the generalization and consistency of the model, as well as similarities and differences between NF learning strategies across participants. Our results suggest that future NF research could exploit distributed information in the brain to improve the efficiency of the NF signal [18], [83], [84], or to monitor and even guide the control strategy used by the participants. Finally, this approach is not limited to analyzing data from NF experiments, but can in principle be useful for gaining new insights in other types of longitudinal data from learning experiments.

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TABLE I

Group 1 ranking	t	β	anatomical location	Group 2 ranking	t	β	anatomical location
1	-6.115	-0.030	Right Middle Occipital Gyrus, Superior Occipital Gyrus	1	4.815	0.012	Midcingulate Cortex
2	-2.889	-0.024	Right Angular Gyrus, Middle Occipital Gyrus	2	-3.316	-0.018	Right Angular Gyrus
3	-2.888	-0.023	Right Insula	3	2.860	0.030	Right Middle Frontal Gyrus
4	2.796	0.012	Left Crus I	4	-2.725	-0.063	Right Inferior Parietal Lobule
5	-2.764	-0.022	Right Caudate	5	-2.650	-0.029	Left Inferior Frontal Gyrus, Orbitofrontal Gyrus
6	-2.753	-0.015	Right Inferior Frontal Gyrus	6	2.589	0.044	Right Supramarginal Gyrus, Inferior Parietal Gyrus
7	-2.738	-0.006	Right Thalamus Caudate, Putamen	7	-2.283	-0.025	Left inferior Temporal Gyrus, Middle Temporal Gyrus
8	-2.625	-0.026	Left Crus I, Crus II, Lobule VI	8	2.146	0.037	Right Superior Temporal Gyrus
9	2.601	0.021	Right Middle Frontal Gyrus	9	-1.925	-0.018	Right Supplementary Motor Area
10	-2.559	-0.032	Right Midcingulate Cortex	10	1.888	0.018	Left Middle Temporal Gyrus
11	-2.408	-0.015	Left Frontal Operculum, Inferior Frontal Gyrus	11	-1.887	-0.029	Left Retrosplenial Cortex, Posterior Cingulate Cortex
12	-2.082	-0.032	Left Precentral Gyrus, Postcentral Gyrus	12	-1.828	-0.037	Left Angular Gyrus
13	-2.033	-0.016	Left Parahippocampal Gyrus	13	-1.803	-0.051	Right Middle Occipital Gyrus, Superior Occipital Gyrus
14	1.897	0.010	Left Thalamus	14	1.609	0.028	Left Middle Temporal Gyrus, Angular Gyrus
15	-1.888	-0.019	Right Superior Frontal Gyrus	15	1.446	0.010	Midcingulate Cortex, Posterior Cingulate Cortex
16	1.809	0.016	Right Inferior Parietal Gyrus, Supramarginal Gyrus, Angular Gyrus	16	-1.385	-0.029	Left Middle Frontal Gyrus, Superior Frontal Gyrus, Precentral Gyrus
17	1.761	0.019	Left Supramarginal Gyrus, Inferior Parietal Gyrus	17	1.382	0.035	Right Retrosplenial Cortex, Posterior Cingulate Cortex
18	1.695	0.039	Right Superior Frontal Gyrus, Middle Frontal Gyrus	18	1.342	0.038	Left Superior Parietal Gyrus, Inferior Parietal Gyrus, Precuneus, Angular Gyrus
19	-1.689	-0.024	Right Superior Frontal Gyrus	19	1.321	0.020	Right Inferior Parietal Gyrus, Supramarginal Gyrus, Angular Gyrus
20	1.611	0.041	Left Middle Frontal Gyrus, Superior Frontal Gyrus, Precentral Gyrus	20	-1.296	-0.040	Left Superior Temporal Gyrus Heschl's Gyrus
21	1.607	0.019	Right Frontal Operculum, Inferior Frontal Gyrus	21	-1.279	-0.025	Left Supramarginal Gyrus, Inferior Parietal Gyrus
22	1.595	0.012	Right Lobule VI, Crus I	22	-1.269	-0.021	Right Supramarginal Gyrus, Superior Temporal Gyrus, Middle Temporal Gyrus
23	-1.551	-0.020	Precuneus	23	1.253	0.018	Left Middle Occipital Gyrus
24	-1.498	-0.015	Right Angular Gyrus	24	1.206	0.026	Calcarine Sulcus
25	1.493	0.037	Left Angular Gyrus	25	1.189	0.005	Left Thalamus
26	1.470	0.030	Left Middle Frontal Gyrus, Superior Frontal Gyrus	26	1.168	0.051	Right Middle Frontal Gyrus, Right Superior Frontal Gyrus
27	-1.463	-0.041	Right Superior Parietal Gyrus, Precuneus	27	1.149	0.015	Left Middle Frontal Gyrus, Superior Frontal Gyrus
28	1.430	0.015	Right Retrosplenial Cortex, Posterior Cingulate Cortex	28	-1.129	-0.009	Left Inferior Frontal Gyrus
29	-1.419	-0.037	Right Superior Frontal Gyrus	29	-1.069	-0.012	Right Thalamus Caudate, Putamen
30	1.414	0.010	Inferior Frontal Gyrus	30	1.057	0.014	Left Inferior Parietal Sulcus
31	1.358	0.023	Precuneus	31	-1.056	-0.013	Left Hippocampus
32	1.344	0.003	Pons	32	1.049	0.020	Right Middle Frontal Gyrus
33	-1.324	-0.020	Right Crus I	33	-1.048	-0.026	Right Frontal Operculum, Inferior Frontal Gyrus
34	-1.286	-0.018	Right Middle Frontal Gyrus, Right Superior Frontal Gyrus	34	-1.046	-0.007	Left Thalamus
35	1.282	0.015	Left Middle Occipital Gyrus	35	1.045	0.007	Right Hippocampus
36	1.171	0.015	Anterior Cingulate Cortex, Medial Prefrontal Cortex, Supplementary Motor Area	36	1.026	0.008	Inferior Frontal Gyrus
37	1.089	0.007	Right Supramarginal Gyrus, Superior Temporal Gyrus, Middle Temporal Gyrus	37	-1.025	-0.016	Right Superior Frontal Gyrus, Middle Frontal Gyrus
38	-1.087	-0.001	Left Middle Temporal Gyrus	38	0.979	0.022	Left Frontal Operculum, Inferior Frontal Gyrus
39	-1.085	-0.013	Left Middle Temporal Gyrus, Superior Temporal Gyrus, Supramarginal Gyrus, Angular Gyrus	39	-0.961	-0.007	Right Parahippocampal Gyrus
40	-1.039	-0.019	Left Inferior Parietal Sulcus	40	-0.900	-0.010	Right Insula
41	-1.020	-0.015	Right Middle Frontal Gyrus	41	0.888	0.010	Left Precuneus
42	0.949	0.012	Right Lobule VI, Crus I	42	0.885	0.011	Right Posterior Insula
43	0.949	0.008	Left Middle Frontal Gyrus	43	-0.882	-0.005	Right Caudate
44	0.919	0.046	Right Supramarginal Gyrus, Inferior Parietal Gyrus	44	0.854	0.006	Right Lobule VIII, Lobule VIIb
45	-0.909	-0.010	Right Thalamus	45	0.827	0.019	Right Superior Parietal Gyrus, Precuneus
46	-0.876	-0.029	Left Superior Parietal Gyrus, Inferior Parietal Gyrus, Precuneus, Angular Gyrus	46	-0.806	-0.022	Medial Prefrontal Cortex, Anterior Cingulate Cortex, Orbitofrontal Cortex
47	-0.848	-0.006	Midcingulate Cortex, Posterior Cingulate Cortex	47	-0.806	-0.020	Precuneus
48	0.834	0.010	Left and Right Thalamus	48	0.804	0.011	Left Angular Gyrus
49	0.823	0.018	Left Superior Temporal Gyrus Heschl's Gyrus	49	0.801	0.012	Left Middle Frontal Gyrus
50	0.789	0.003	Left Inferior Frontal Gyrus	50	0.764	0.013	Right Thalamus
51	-0.764	-0.007	Left inferior Temporal Gyrus, Middle Temporal Gyrus	51	0.728	0.004	Right Thalamus
52	0.763	0.033	Right Supplementary Motor Area	52	0.722	0.006	Right Midcingulate Cortex
53	-0.761	-0.014	Right Middle Frontal Gyrus	53	-0.674	0.002	Right Middle Frontal Gyrus
54	0.761	0.012	Right Angular Gyrus	54	0.671	0.002	Right Lobule VI, Crus I
55	0.753	0.004	Right Superior Temporal Gyrus	55	-0.627	-0.004	Lobule VI
56	-0.732	-0.011	Right Middle Temporal Gyrus	56	0.602	0.009	Left Middle Occipital Gyrus, Superior Occipital Gyrus
57	0.700	0.001	Left Angular Gyrus	57	0.588	0.006	Left Posterior Insula, Putamen
58	0.692	0.004	Right Thalamus	58	-0.584	-0.005	Left Crus I
59	0.664	0.030	Left Retrosplenial Cortex, Posterior Cingulate Cortex	59	-0.545	-0.003	Left Middle Frontal Gyrus,
60	-0.649	-0.006	Right Lobule VIII, Lobule VIIb	60	0.539	0.008	Left Precentral Gyrus, Postcentral Gyrus
61	-0.645	-0.020	Medial Prefrontal Cortex, Anterior Cingulate Cortex, Orbitofrontal Cortex	61	-0.532	-0.001	Right Lobule IX
62	0.586	0.006	Calcarine Sulcus	62	0.506	0.004	Right Lobule VI, Crus I
63	0.558	0.008	Right Thalamus	63	-0.498	-0.005	Right Precentral Gyrus, Postcentral Gyrus
64	-0.532	-0.017	Left Thalamus	64	-0.476	-0.007	Lobule VI
65	-0.522	-0.005	Left Middle Frontal Gyrus	65	0.474	0.004	Right Inferior Frontal Gyrus
66	-0.436	-0.001	Left Thalamus	66	-0.466	-0.010	Left Parahippocampal Gyrus
67	0.434	0.007	Right Parahippocampal Gyrus	67	-0.447	-0.010	Right Crus I
68	0.427	0.014	Left Thalamus, Caudate	68	-0.437	-0.002	Left Lobule VI, Crus I
69	-0.422	-0.010	Right Hippocampus	69	0.401	0.006	Right Angular Gyrus, Middle Occipital Gyrus
70	0.392	0.003	Left Inferior Temporal Gyrus	70	0.396	0.004	Precuneus
71	-0.371	0.000	Right Posterior Insula	71	-0.388	-0.007	Right Angular Gyrus
72	0.358	0.009	Right Inferior Parietal Lobule	72	-0.370	-0.002	Bilateral Lobule IV, Lobule V, Lobule VI
73	0.357	0.001	Left Middle Occipital Gyrus, Superior Occipital Gyrus	73	-0.341	-0.001	Left Crus I, Crus II, Lobule VI
74	-0.348	-0.005	Left Precuneus	74	0.338	0.000	Right Superior Frontal Gyrus
75	-0.332	-0.002	Left Lobule VIII, Lobule VIIb	75	-0.332	-0.016	Anterior Cingulate Cortex, Medial Prefrontal Cortex, Supplementary Motor Area
76	-0.276	-0.005	Left Middle Frontal Gyrus	76	-0.318	-0.001	Left Inferior Temporal Gyrus
77	0.271	0.001	Right Lobule IX	77	-0.314	-0.004	Left Middle Frontal Gyrus
78	-0.233	0.000	Left Thalamus	78	0.312	0.007	Right Inferior Frontal Gyrus
79	0.233	-0.002	Lobule VI	79	0.310	0.002	Left Thalamus
80	0.227	0.001	Left Insula	80	0.308	0.004	Left Thalamus
81	-0.214	-0.003	Left Middle Temporal Gyrus, Angular Gyrus	81	0.295	0.009	Left Insula
82	0.206	0.002	Left Lobule VI, Crus I	82	0.260	0.007	Right Superior Frontal Gyrus
83	-0.187	0.004	Bilateral Lobule IV, Lobule V, Lobule VI	83	-0.245	-0.005	Right Superior Frontal Gyrus
84	-0.182	0.000	Left Hippocampus	84	0.241	-0.002	Right Middle Temporal Gyrus
85	0.177	-0.002	Right Inferior Frontal Gyrus	85	0.240	-0.003	Left Thalamus, Caudate
86	-0.169	-0.002	Midcingulate Cortex	86	0.201	0.006	Left and Right Thalamus
87	0.162	-0.009	Left Posterior Insula, Putamen	87	-0.172	-0.003	Left Middle Temporal Gyrus, Superior Temporal Gyrus, Supramarginal Gyrus, Angular Gyrus
88	0.155	0.005	Right Precentral Gyrus, Postcentral Gyrus	88	0.102	0.001	Pons
89	-0.134	0.003	Lobule VI	89	0.034	-0.002	Left Lobule VIII, Lobule VIIb
90	0.042	0.006	Left Inferior Frontal Gyrus, Orbitofrontal Gyrus	90	-0.019	-0.005	Right Thalamus