Estimations of the weather effects on brain functions using functional MRI – a cautionary tale

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Abstract

The influences of environmental factors such as weather on human brain are still largely unknown. A few neuroimaging studies have demonstrated seasonal effects, but were limited by their cross-sectional design or sample sizes. Most importantly, the stability of MRI scanner hasn't been taken into account, which may also be affected by environments. In the current study, we analyzed longitudinal resting-state functional MRI (fMRI) data from eight individuals, where the participants were scanned over months to years. We applied machine learning regression to use different resting-state parameters, including amplitude of low-frequency fluctuations (ALFF), regional homogeneity (ReHo), and functional connectivity matrix, to predict different weather and environmental parameters. For a careful control, the raw EPI and the anatomical images were also used in the prediction analysis. We first found that daylight length and temperatures could be reliability predicted using cross-validation using resting-state parameters. However, similar prediction accuracies could also achieved by using one frame of EPI image, and even higher accuracies could be achieved by using segmented or even the raw anatomical images. Finally, we verified that the signals outside of the brain in the anatomical images and signals in phantom scans could also achieve higher prediction accuracies, suggesting that the predictability may be due to the baseline signals of the MRI scanner. After all, we did not identify detectable influences of weather on brain functions other than the influences on the stability of MRI scanners. The results highlight the difficulty of studying long term effects on brain using MRI.

Keywords: daylight length, environmental effects on brain, machine learning regression, resting-state, scanner stability, temperature, weather.

1. Introduction

Daily environmental factors such as weather and seasonality affect mood and cognitive functions (Keller et al. 2005; Denissen et al. 2008; Cedeño Laurent et al. 2018; IJzerman et al. 2018; Lim et al. 2018), and may lead to pathological affective disorder (Kurlansik and Ibay 2012). The effects on individuals may be small, but the collective effects may lead to broader impacts, e.g. on stock markets (Saunders 1993; Hirshleifer and Shumway 2003). To better understand the effects of weather and seasonality on mood or cognitions, it is critical to study their effects on brain functions. A few human neuroimaging studies have explored this association. Seasonal effects on brain functions as measured by functional MRI (fMRI) have been observed both in resting-state (Choe et al. 2015) and when performing cognitive tasks (Meyer et al. 2016). Some neural transmitter activity in the striatum also showed seasonal effects, i.e. serotonin transmitter binding as measured by ^{11}C–labeled 3-amino-4-(2-dimethylaminomethyl-phenylsulfanyl)-benzonitrile ($[^{11}$C]DASB) positron emission tomography (PET) (Praschak-Rieder et al. 2008; Kalbitzer et al. 2010; Mc Mahon et al. 2016) and dopamine synthesis as measured by ^{18}F-DOPA PET (Eisenberg et al. 2010; Kaasinen et al. 2012). A study even reported seasonal changes of hippocampal volumes in human subjects (Miller et al. 2015).

There are several limitations in these neuroimaging studies. First, most of these studies are cross-sectional, which is limited by the large individual differences in brain functions (Gordon et al. 2017). In addition, most of the studies examined roughly defined seasonal effects or yearly periodical effects. But the exact phase of the seasonal variations may be different from the roughly defined four seasons. Sometimes the yearly effects showed different phases (Meyer et al. 2016), suggesting more complicated relationships of environmental factors on brain functions. Therefore, it is critical to examine which environmental parameters such as weather have more contributions to the seasonal effects. Among different weather parameters, daylight length and temperature represent the significant environmental differences in seasonal fluctuations. Gillihan et al has explored the weather effects on brain functions using a small cross-sectional sample (Gillihan et al. 2011). They identified a weather index related to

mood, and showed that the weather index was correlated with resting-state cerebral blood flow as

measured by arterial spin labeled (ASL) perfusion fMRI mainly in the insula. But more systematic

examinations of weather effects on brain functions have not been performed. Lastly, the most commonly

used neuroimaging method is blood-oxygen-level dependent (BOLD) signal based fMRI, where the

interpretation of the results should consider neuronal level, neurophysiological level, and the underlying

physical level of the scanner. Specifically, if some effects on fMRI signals were observed, they may be

due to the changes of neuronal activity, which is favorable to psychologists and psychiatrist. But the

effects may also due to the changes of neurovascular coupling (Di et al. 2013; Yuan et al. 2013), changes

of brain structures, or even the stability of the MRI scanner. Therefore, when examining the weather

effects on brain functions, the alternative factors need to be considered and carefully controlled.

The purpose of the current study is to estimate to what extent resting-state brain functions were

affected by weather. We analyzed longitudinal resting-state fMRI data from eight individuals from three

datasets, where the individuals were scanned over periods of months to years (Choe et al. 2015; Poldrack

et al. 2015; Filevich et al. 2017). One challenge for estimating weather effects is that the effects may be

small. Therefore, we applied a machine learning regression approach to evaluate the amount of effects.

Because multiple brain regions have been implicated in seasonal effects, e.g. basal ganglia (Praschak-

Rieder et al. 2008; Kalbitzer et al. 2010; Mc Mahon et al. 2016), insula (Gillihan et al. 2011), and

hippocampus (Miller et al. 2015), small regional effects may be aggregated into detectable effects using

machine learning technique. We performed within-subject prediction analysis at single subject level. We

asked what weather parameters have the most effects on resting-state brain functions, which can be

represented as high prediction accuracies in predictions of these parameters. In order to rule out possible

confounding effects that might give rise to prediction, we also performed control prediction analyses.

First, we analyzed anatomical MRI images to check whether the observed prediction could be attributed

to anatomical variations. Second, we checked images from phantom data to examine whether the

prediction could be attributed to the stability of the MRI scanner.

2. Materials and methods:

2.1. MRI datasets

Several multi-session resting-state fMRI datasets were pooled together, where the subjects were scanned

over periods of months to years. The first subject was derived from the Kirby sample (Choe et al. 2015),

where the single subject was scanned for 156 sessions over three and half years. The second subject was

from the Myconnectome sample (Poldrack et al. 2015), where the subject was scanned 90 times over one

and half years. The remaining six subjects were from the Day2day sample (Filevich et al. 2017), where

the subjects were scanned over a similar span of about half a year. The detailed subject and scan

information is listed in Table 1.

	Dataset	Sex	Age	# of sessions	First scan	Last scan	# of volumes	TR (s)	Voxel size (mm^3)
1	Kirby	M	40	156	2009/12/07	2013/06/20	198	2	3 x 3 x 4
2	MyConnectome	M	45	83	2012/10/23	2014/03/11	500	1.16	2.4 x 2.4 x 2.4
3	Day2day	F	23	50	2013/07/03	2013/12/18	148	2	3 x 3 x 3.6
4	Day2day	F	31	48	2013/07/03	2014/01/08	148	2	3 x 3 x 3.6
5	Day2day	F	29	45	2013/07/03	2014/01/27	148	2	3 x 3 x 3.6
6	Day2day	F	24	46	2013/07/02	2013/12/19	148	2	3 x 3 x 3.6
7	Day2day	M	30	39	2013/07/09	2014/02/12	148	2	3 x 3 x 3.6
8	Day2day	F	29	48	2013/07/03	2014/02/20	148	2	3 x 3 x 3.6

Table 1 Subject and scan information for the eight subjects with longitudinal scans. The number of

sessions represents the effective number of sessions after dropout due to missing data or large head

motions. The numbers of volumes represent the number of volumes used in the analysis after removing

the first several volumes.

The MRI data from the Kirby sample were scanned using a 3T Philips Achieva scanner. The data

from the Myconnectome sample were scanned using a 3T Siemens Skyra scanner using a 32-channel

head coil. And the data from the Day2day project were scanned using a 3T Siemens Magnetom Trio

scanner using a 12-channel head coil. For each subject, resting-state fMRI data with multiple sessions

were available. Within a subject, the resting-state fMRI were scanned using the same imaging parameters,

but the parameters varied between different sites. Some essential resting-state fMRI parameters are listed

in Table 1. For more details, we refer the readers to the original articles.

High resolution anatomical MRI images were available for only a few sessions in the Kirby and

Myconnectome datasets for each subject. A MRI image of one session was used to register all the

functional images to standard Montreal Neurological Institute (MNI) space. For the Day2day dataset,

structural MRI images were available for all the sessions. Only the structural MRI image of the last

session of each subject was used to aid preprocessing of the fMRI images. All the structural images of

the Day2day project were also used in the control prediction analysis.

Lastly, we obtained MRI scanner quality assurance agar phantom data from the Day2day site.

The images were scanned between June 2013 and February 2014 with a weekly basis (37 sessions in

total). The data were acquired using gradient echo (GRE) sequence with the same coil as the one used for

acquisition of the human data. Two images were acquired for each session. The parameters include: TR

$= 2000$ ms; TE $= 30$ ms; FOV $= 22$ cm; matrix $= 64 \times 64$; slice number $= 28$; slice thickness $= 4$ mm (1

mm gap).

2.2. Environmental data

The fMRI data were acquired from three different cities in two continents, Baltimore USA (Kirby),

Austin USA (Myconnectome), and Berlin Germany (Day2day), which reflect different types of climates.

The latitudes of these four cities are approximately 39 ^{o}N, 30 ^{o}N, and 52 ^{o}N, respectively. The weather

data for the two US cities were downloaded from (US) National Centers for Environmental Information

website (https://www.ncdc.noaa.gov/cdo-web/). The Local Climatological Data from Maryland Science

Center Station and Austin Camp Mabry Station were used to represent the weather for the Kirby and

Myconnectome datasets, respectively. We used the following measures, maximum and minimum

temperatures, air pressure, wind speed, humidity, and precipitation. For those with missing data, we also

checked Daily Summaries data from the NOAA website. The weather data for the Day2day dataset were

collected by the German researchers. Daily sunshine hours were not used, because they were not

available for the other datasets.

We also included daylight length in the current analysis. It was already available in the NOAA

Local Climatological Data. For the Day2day data, we calculated the daylight length in Berlin according

to its geographic location through the website of the Astronomical Applications Department of the U.S.

Naval Observatory computes (http://aa.usno.navy.mil/data/docs/Dur_OneYear.php). For the Day2day

dataset, there are also three parameters that reflect local environmental variations, i.e. scanner room

temperature, scanner room humidity, and scanner Helium level. These three parameters were also used in

the prediction analysis when using Day2day data to do prediction.

2.3. MRI data processing

2.3.1. Resting-state fMRI Preprocessing

Data processing and statistical analysis were performed using MATLAB (R2017b). SPM12

(http://www.fil.ion.ucl.ac.uk/spm/ ; RRID:SCR_007037) was used for fMRI data preprocessing. The first

2, 18, and 2 functional images for each session were discarded for Kirby, Myconnectome, and Day2day

datasets, respectively, remaining 198, 500, and 148 images for each session. For each subject, all the

functional images were realigned to the first session. All the prediction analysis was performed in native

space of each subject. The anatomical images were coregistered to the mean functional image, and then

segmented into gray matter (GM), white matter (WM), cerebrospinal fluid (CSF), and other tissues. At

each voxel, Friston's 24 head motion model, the first five principle components from WM signals and the

first five principle components from CSF signals were regressed out, and then a band-pass filtering was

applied at frequency of 0.01 and 0.1 Hz. The images were not spatially smoothed, because there was no

voxel-wise univariate analysis involved.

2.3.2. ALFF, ReHo, and connectivity matrices

We calculated three resting-state parameters to represent resting-state brain functions, i.e. amplitude of

low-frequency fluctuation (ALFF) (Zang et al. 2007) and regional homogeneity (ReHo) (Zang et al. 2004)

to represent regional properties, and connectivity matrix to represent inter-regional connectivity property.

ALFF and ReHo were calculated using REST toolbox (Song et al. 2011). Essentially, ALFF calculated

the power of the time series signals between 0.01 to 0.08 Hz at every voxel, resulting in an ALFF map for

each session. ReHo calculated the correlations of the current voxel with the 26 neighboring voxels, which

also resulted in a ReHo map for each session. The ALFF and ReHo values for each session within the

subject's GM mask were concatenated as a vector for further analysis. The subject specific GM masks

were defined as GM intensity greater than 0.5 based on the segmentation results from the subject's own

anatomical image. Because the GM masks were defined in the native spaces and the fMRI resolution

varies across datasets, the number of within mask voxels also varied (from 20,780 to 55,368).

Correlation matrices were calculated among 164 regions of interests (Dosenbach et al. 2010; Di

and Biswal 2019). Spherical ROIs were first defined in MNI space with a radius of 8 mm, and

transformed into the native space for each subject. There were in total 13,366 connectivity values (164 x

(164 - 1) / 2), which were concatenated as a vector for the prediction analysis. The correlation values

were transformed into Fishers' z scores.

2.3.3. Head motion and other potential confounding variables

To minimize the confounding of head motion in the prediction analysis, we first removed sessions with

large head motions. We calculated frame-wise displacement in translation and rotation directions (Di and

Biswal 2015). A session's data with maximum frame-wise displacement greater than 1 mm or 1° were

discarded. No sessions were removed in the Kirby data, and seven sessions (7.8%) were removed for the

Myconnectome data. In the Day2day dataset, at most two sessions were removed for each subject.

Secondly, we regressed out 24 motion variables using Friston's head motion model (Friston et al. 1996),

which has been shown to be effective to minimize the effects of head motion on resting-state measures

(Yan, Cheung, et al. 2013). Lastly, mean frame-wise displacement of both directions were regressed out

from a predicted environmental variable before it was entered into the prediction analysis.

2.3.4. Structural MRI processing

For the Day2day dataset, the MPRAGE anatomical MRI images were available for all the sessions.

Therefore, we used the anatomical images as a control condition for weather prediction. The analysis was

also performed in subject's native space. The anatomical images from all the sessions of a subject were

realigned and resliced to the image of the first session. Then each session's image was segmented

separately, and the segmented tissue probability maps of GM, WM, and CSF were obtained in the native

space. We defined GM, WM, and CSF masks as averaged probability greater than 0.5 in respective tissue

type. GM, WM, and CSF probability in their masks were extracted, respectively, to be used in the

prediction analysis.

2.3.5. Phantom image processing

For each session, the two images were realigned, and an averaged image was calculated. Because the

phantom was placed in the similar location, no cross-session registration was performed. We first

calculated correlations between daylight length and image values in every voxel, resulting in a correlation

image. Next, a cubic mask in the center of the image was defined. The signals within the mask were

extracted for the prediction analysis.

2.4. Prediction analysis

2.4.1. Prediction analysis scheme

The goal of the analysis is to estimate the prediction values of resting-state parameters on different

weather or meteorological parameters. The analysis was performed for each of the resting-state

parameters to predict each of the seven weather parameters. And we asked which weather parameters

could be better predicted by which resting-state parameters. The prediction analysis was all done in a

within-subject manner. Cross-validation was used to evaluate the prediction accuracy.

In addition to use these resting-state parameters, we also performed a series of control analysis to

use other potential confounding parameters to predict the environmental parameters. First, we used the

first fMRI image of each session after realignment to perform prediction analysis. Although the single

image still reflects BOLD effects, it may be more contributed by brain structural variations. Secondly, to

future rule out the structural contribution, we used segmented tissue probabilities of GM, WM, and CSF

from their respective tissue masks to perform prediction analysis. Thirdly, we also extracted raw image

values from the MPRAGE images in the three masks to serve as another control conditions. In addition, a

cuboid mask was also defined for each subject, which was located outside the brain. The raw image

values from the MPRAGE images from the air mask was used to control for baseline MRI signals.

Finally, since all of the above mentioned analyses indicated prediction values to predict environmental

parameters, especially daylight length, we further analyzed the quality assurance phantom data, and used

the signals in the agar phantom area to perform prediction analysis to predict daylight length.

2.4.2. Machine learning regression analysis

We used linear machine learning regression model to perform prediction analysis. The general form of

the prediction model is a linear regression model as the following:

$$y = X \cdot \beta + \varepsilon$$

where y is a *1 x n* vector of a predicted weather parameter, X is a *n x m* matrix of a resting-state parameter,

β is the model parameters, and ε is the residual. N represents the number of observations, which in the

current analysis were the number of sessions for the longitudinal data and the number of subjects for the

cross-subject data. M represents the number of prediction variables, which could be the number of voxels

in the ALFF or ReHo maps (see Table 1) or the number of connections (13,366) in the connectivity

matrices. Here, m is much larger than n. Therefore, we used ridge regression to estimate the β

parameters. Briefly speaking, instead of trying to achieve the goal of minimizing the sum of square

means of the model prediction:

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$$\min_{\beta} \sum (y - X \cdot \beta)^2$$

236   Ridge regression adds one more regularized term:

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$$\min_{\beta} \sum (y - X \cdot \beta)^2 + \lambda \cdot \sum \|\beta\|^2$$

where λ represents the regularization parameter. The regularization term can constrain the sizes of beta

values, thus preventing over fitting of the model. In the current analysis, we used the MATLAB function

fitrlinear to perform the prediction analysis. There are other methods available, such as LASSO and

elastic net, but a recent study suggested that ridge regression and elastic net can yield similar prediction

accuracies while LASSO might perform worse in the scenario that the number of observations is much

smaller than the number of features (Cui and Gong 2018).

There are three steps in the prediction analysis, 1) tuning regularization parameter λ to find the

optimal λ (λ tuning), 2) training the model using the training dataset and the optimal λ to obtain a

prediction model β (model training), and 3) estimating prediction accuracy by calculating correlations

between predicted and actual values in a separate testing sample (cross-validation). Cross-validation was

used to make sure that the estimated prediction accuracy was independent from the training data.

Therefore, it is important for the within-subject and cross-subject prediction. For the between-subject

prediction, however, prediction accuracy was estimated from a separate subject; therefore no cross-

validation was required.

Because of the limited number of data in one fold (13 observations in the least case), 3-fold cross-

validation was adopted. We used a nested tuning strategy to optimize the parameter λ (Figure 1A) (Cui

and Gong 2018). Specifically, we first used 3-fold cross-validation to hold out one third of the data as an

independent testing dataset, and used the remaining two thirds of the data as training and parameter

tuning dataset. The data were first sorted according to the tested weather parameter, and the three folds

were defined as the 1st, 4th, 7th, …, 2nd, 5th, 8th, …, and 3rd, 6th, 9th, … sessions of the data, respectively.

Within the two thirds training and parameter tuning dataset, we first performed a nested loop of 3-fold

analysis. Specifically, one third of the data were hold out, and the remaining of the two thirds of data

were used to train the regression model using a set of λ values, from 10^{-5} to 10^{-1} in the logarithmic scale

with a total 15 values. The one third of inner loop testing data was used to test the accuracy of the

prediction by calculating the correlation between predicted and actual weather parameter values. This

procedure was performed for three times for the three folds, and the mean accuracies were calculated for

each of the λ values. The λ value with the highest mean accuracy was used for the outer layer training

data to train the model. The model was then applied to the outer layer testing data to estimate prediction

accuracies. The three accuracy values from the 3 folds were averaged to represent an estimate of

accuracy for a subject.

3. Results

3.1. Weather predictions using the resting-state data

We first performed prediction analysis on different weather parameters using the ALFF maps, ReHo

maps, connectivity matrices, as well as single frame of EPI images as a control condition (Figure 1). In

general daylight length (DayL) and maximum and minimum environmental temperatures (T max and T

min) had prediction accuracies. And daylight length usually had the highest prediction accuracies. All

other environmental parameters had very low prediction accuracies. In terms of resting-state parameters,

the ALFF map usually had the highest prediction value. The averaged prediction accuracy of daylight

length was 0.42. However, using a single frame of EPI images could achieve comparable prediction

accuracies as the ALFF maps.

Figure 1 Prediction accuracies (correlations) of the ALFF (amplitude of low-frequency fluctuations)

maps, ReHo (regional homogeneity) maps, and connectivity matrices on different weather parameters.

Each dot represents one subject's mean prediction accuracy. The center white line and inner dark bar

represent the mean and 95% confidence interval, respectively, and the outer light bar represents one

standard deviation to the mean. One sample t test was performed for each condition, and the red * on the

top represents statistical significant cross-subject effects at $p < 0.05$ after FDR (false discovery rate)

correction for all the 21 predictions. The figure was made by using notBoxPlot

(https://github.com/raacampbell/notBoxPlot).

It is still a question that whether the weather predictions using the resting-state parameters and

single EPI are based on similar or different information. Since the daylight length had the highest

prediction accuracies, we focused on its prediction. We combined ALFF with EPI or ReHo with EPI to

predict daylight length to check whether combining the two modalities can boost the prediction accuracies.

Unfortunately, combing two modalities yielded very similar prediction accuracies as those using single

EPI images or ALFF images (Figure 2). Therefore, ALFF and ReHo did not convey more information

than a single frame EPI image to predict daylight length.

Figure 2 Prediction accuracies to daylight length using the ALFF, Reho, raw EPI maps, and the

combination of EPI with ALFF or EPI.

3.2. Prediction analysis using the anatomical images

If single frame EPI images can predict weather parameter like daylight length, then the question becomes

whether the prediction is due to brain functional activity, structural information, or other factors. We

therefore performed the same prediction analysis using anatomical images, where were available in the

six subject in the Day2day dataset. We first performed prediction analysis using segmented GM, WM, or

CSF images using their respective tissue mask (Figure 3). The results showed very similar prediction

patterns for different environmental parameters as using resting-state parameters. That is, the daylight

length and environmental temperatures had the highest prediction accuracies. The prediction accuracies

using all the three tissue probability maps were above 0.5, which were higher than using any resting-state

parameters. However, what more interesting was that even higher prediction accuracies could be

achieved using raw anatomical images in different tissue masks. The prediction accuracies were higher

than 0.6 when using raw anatomical image signals in the GM or CSF masks. Finally, we defined a

cuboid mask outside the brain (see Figure 4A as an example), and used the raw MRI signals in the mask

to perform prediction analysis. Surprisingly, the analysis also showed similar pattern of prediction

314 accuracies. The prediction accuracy on daylight length using the air mask was 0.47, which was lower

315 than using all other anatomical parameters but still higher than using any of the resting-state parameters.

317   **Figure 3** Cross-subject predictions of weather parameters using the amplitude of low-frequency

318   fluctuations (ALFF) maps (top row), regional homogeneity (ReHo) maps (middle row), and connectivity

319   matrices (bottom row) from the six subjects of the Day2day dataset. T max and T min represent daily

320   maximum and minimum temperature, respectively.

To further explore the baseline MRI signals conveyed in the air mask, we calculated correlations

between MPRAGE signals and daylight length in all the voxels in the air mask in all the six subjects

(Figure 4B). There were small global effects of correlations between MPRAGE signals and daylight

length. The global effects of correlations were strongly correlated with the prediction accuracies across

subjects (Figure 4C), indicating that the global correlation is the driving information that gave rise to the

prediction accuracy.

Figure 4 A) An example of the air mask from one subject overlaid to the subject's mean anatomical

image. B) Histograms of the correlations between the MRI signals and daylight length of all the voxels in

the air mask. Each line with different colors represents one subject. C) There is an extremely high

negative correlation between the mean correlations in the air mask and the prediction accuracies of using

the air mask voxels to predict daylight length.

3.3. Control analysis using the phantom images

To further confirm the baseline signal changes, we analyzed the weekly quality control phantom data

around the same period of the Day2day project. We first calculated voxel-wise correlations between the

MRI signal and daylight length (Figure 5A). It clearly showed that in the phantom region, there were

high negative correlations. We defined a cubic mask in the center of the image, and the distribution of

correlations of all the voxels in the mask were plotted in Figure 5B. The mean and median correlation in

the cubic mask was *-0.75* and *-0.77*, respectively. Lastly, we performed the same kind of prediction

analysis using all the voxels in the mask to predict daylight length, and the cross-validated mean accuracy

was 0.72.

Figure 5 A) Voxel-wise Correlation maps between MRI image values and daylight length. The dash-line

contour indicates the location of the cubic mask. B) Histogram of the correlations between MRI image

values and daylight length in all the voxels in the cubic mask.

4. Discussion

By applying machine learning regression to longitudinal fMRI data that were scanned over months to

years, we demonstrated that we can predict environmental parameters, especially daylight length and

temperature, by using resting-state fMRI parameters. However, a series of control analysis showed that

using a single EPI image, segmented tissue images, and raw MPRAGE image signals in different tissue

masks and even in a mask outside the brain could all predict the environmental parameters. The resting-

state parameters did not add prediction values to single frame EPI images. These results indicated the

prediction of weather parameters, especially daylight length, cannot be explained as the weather effects

on brain functions. Rather, the prediction may reflect MRI scanner baseline signal variations that were

affected by environmental parameters. The analysis of the quality control phantom images supported our

speculation.

Among all the environmental parameters analyzed the daylight length had the highest prediction

accuracies. It's noteworthy that the prediction accuracies of daylight length were even higher than what

in the predictions of MRI scanner room temperature and humidity. Therefore, there may be some other

local factors that mediate the effects of daylight length on scanner stability. A study has shown that the

gaseous oxygen level in the magnet field can influence the MRI signals (Bates et al. 1995). It is

reasonable to speculate that the oxygen level in the scanner room may fluctuate across seasons due to

different ventilation conditions. In addition, the cooling systems of the scanner may be affected by either

electricity supply stability or cooling water temperature. Given that the MRI is such a sophisticated

machine, there may be other factors that mediate the association between daylight length and scanner

stability.

The current results highlighted the difficulty to study long term effects such as weather on brain

structures and functions using MRI. Consistent with two previous fMRI studies (Choe et al. 2015; Meyer

et al. 2016), we did find weather effects on fMRI measures. But we demonstrated that the weather effects

are likely due to the variations of MRI scanner stability. It is reasonable to speculate that MRI scanner

stability might contribute to the reported seasonal effects (Choe et al. 2015; Meyer et al. 2016). We also

showed that tissue probability measures of GM volumes may also be affected by the scanner stability, so

brain volumetric measures may also be affected by the scanner stability (Miller et al. 2015). The problem

may arise from the fact that most of the fMRI/MRI measures are relative measures. If absolute measures

can be used, e.g. using perfusion measures from arterial spin labeling (ASL) (Detre et al. 1992, 2012),

then the effects from scanner stability may be minimized. In addition, adjusting global signals maybe

helpful to minimize the scanner stability issue (Peelle et al. 2012; Yan, Craddock, et al. 2013). But at the

same time, the global signal was tightly coupled with neural activity (Scholvinck et al. 2010), and

adjusting global signals may cause statistical artifacts (Murphy et al. 2009; Saad et al. 2012).

The current analysis demonstrated that machine learning is a powerful method to be able to pick

up small effects. The phantom data showed the correlations between baseline MRI signals and daylight

length were about 0.7. Although the direct correlations of MRI signals and daylight length when

scanning human participants were blurred (Figure 4A), we could still achieve the similar level as the

prediction accuracies by using machine learning and cross-validation (Figure 3). Indeed, the cross-

validated prediction accuracies were between 0.6 and 0.7, which were very close to the correlation in the

phantom data. Machine learning methods have become more and more popular in studying brain-

behavior relationships (Finn et al. 2015; Cui and Gong 2018) and different mental disorders (Whelan et al.

2014). It is powerful on one hand, but on the other hand, careful control conditions are needed to rule out

possible confounding artifacts.

394   In conclusion, by applying machining learning on resting-state fMRI or structural MRI data, we

can predict several weather parameters, with the highest prediction accuracies to daylight length.

However, the predictions were not likely due to the weather effects on brain functions or structures, but

may due to the weather effects on the stability of MRI scanners. The data highlight the difficulty to use

fMRI/MRI data to study long term effects. And call for cautions to control for scanner stability when

studying long term effects.

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