Default mode network anatomy and function is linked to pediatric concussion recovery

Kartik K. Iyer1, Andrew Zalesky2, Karen M. Barlow1,3,4,5,* & Luca Cocchi6,*

1Child Health Research Centre, Faculty of Medicine, The University of Queensland, Brisbane, Queensland, Australia
2Melbourne Neuropsychiatry Centre & Department of Biomedical Engineering, The University of Melbourne, Victoria, Australia
3Department of Neurology, Queensland Children’s Hospital, Brisbane, Queensland, Australia
4Alberta Children’s Hospital Research Institute, Calgary, Canada
5University of Calgary, Calgary, Canada
6Clinical Brain Networks Group, QIMR Berghofer Medical Research Institute, Herston, Queensland, Australia

Abstract

Objective: To determine whether anatomical and functional brain features relate to key persistent post–concussion symptoms (PPCS) in children recovering from mild traumatic brain injuries (mTBI), and whether such brain indices can predict individual recovery from PPCS. Methods: One hundred and ten children with mixed recovery following mTBI were seen at the concussion clinic at Neurology department Alberta Children’s Hospital. The primary outcome was the Post–Concussion Symptom Inventory (PCSI, parent proxy). Sleep disturbance scores (PCSI subdomain) and the Neurocognition Index (CNS Vital Signs) were also measured longitudinally. PPCS was assessed at 4 weeks postinjury and 8–10 weeks postinjury. Gray matter volumes were assessed using magnetic resonance imaging (MRI) and voxel-based morphometry at 4 weeks postinjury. Functional connectivity was estimated at the same timepoint using resting-state MRI. Two complementary machine learning methods were used to assess if the combination of gray matter and functional connectivity indices carried meaningful prognostic information. Results: Higher scores on a composite index of sleep disturbance, including fatigue, were associated with converging decreases in gray matter volume and local functional connectivity in two key nodes of the default mode network: the posterior cingulate cortex and the medial prefrontal cortex. Sleep-related disturbances also significantly correlated with reductions in functional connectivity between these brain regions. The combination of structural and functional brain indices associated to individual variations in the default mode network accurately predicted clinical outcomes at follow-up (area under the curve = 0.86). Interpretation: These results highlight that the function–structure profile of core default mode regions underpins sleep-related problems following mTBI and carries meaningful prognostic information for pediatric concussion recovery.

Introduction

Incidence rates of traumatic brain injury (TBI) in pediatric populations are on the rise, with median estimates suggesting 691 injuries per 100,000 emergency admissions.1 Most of these injuries are mild TBIs (~90%, mTBI). A large proportion of children have persistent post–concussion symptoms (PPCS) following mTBI2,3 which is defined as the presence of at least two or more post–concussion symptoms that persist for 4 weeks or longer.4 Around 12% of children sustaining an mTBI have PPCS three months following injury.5 There are several PPCS phenotypes with the commonest symptoms being headaches, fatigue, sleep disturbances and cognitive difficulties.6 These symptoms have a detrimental impact on health-related quality of life and long-term outcomes.7–9

Changes in the anatomy and function of whole-brain networks have shown to be associated with abnormalities...
in sleep regulation, fatigue and cognition. Evidence suggests that the default mode network (DMN), which interlinks remote regions such as the posterior cingulate and medial prefrontal cortex, is critically impacted by TBI. Several studies have suggested that the DMN provides a sensitive indicator of the brain’s structural and functional integrity. For example, in stroke and TBI, the anatomy and function of the DMN are compromised and alterations within this network have been linked to poor outcomes. DMN function has indeed been shown to reflect TBI-induced impairments in cognitive flexibility, sleep, and levels of fatigue. In adults, TBI has also been linked to significant changes in the structure of the DMN. In line with these findings, structural changes in white matter pathways connecting DMN regions following mTBI have been linked to corresponding decreases in functional connectivity. DMN functions following TBI have also been attributed to deficits in sustained attention, wherein activity within the DMN is inconsistently regulated. Broadly, functional associations of DMN regions in other systems such as pain, mood and interoceptive awareness, suggest a pervasive role for this network in behaviors commonly encountered in PPCS. Despite these current evidences, the converging impact that mTBI has on the structure and function of pediatric brains is unclear. Moreover, the association between brain changes linked to mTBI, PPCS, and clinical outcomes remains to be proven. On the basis of existing evidence, individual prediction of PPCS recovery may be achievable by examining changes in brain anatomy and function of the DMN following mTBI/concussion.

This report investigates a large prospective sample of children with established PPCS following mTBI. We started by examining whether acute post–injury structural and functional brain changes were linked to PPCS. By adopting an exploratory multivariate analysis, our previous study in the same sample identified key associations between global brain network measures of resting-state functional connectivity and PPCS. In line with existing findings, our previous results showed that a decrease in global DMN resting–state functional connectivity corresponded to increased sleep disturbances and cognitive difficulties in children with mTBI. Here, we expand on our prior findings which suggested that global patterns of functional connectivity (FC) were linked to sleep problems and fatigue in children following mTBI. In this study, we specifically tested the hypothesis that a whole-brain analysis of structure and function, associated to sleep disturbance and fatigue, converges to key regions and connections within the DMN. We employed local anatomical (voxel-based morphometry, VBM) and functional (regional homogeneity, ReHo) measures. These measures were complemented with the analysis of FC between regions showing significant brain–behavior associations. We also posit that these brain indices allow a significant and accurate prediction of clinical and cognitive recovery following mTBI.

**Materials and Methods**

**Subjects**

We included 132 children with a confirmed diagnosis of mTBI: 99 children had PPCS at 4–6 weeks postinjury (henceforth referred as 1-month postinjury) and 33 children who had mTBI at 1-month postinjury recovered. We recruited a convenience sample provided by a sub-study of a clinical trial investigating the efficacy of melatonin in PPCS (PlayGame Trial: NCT01874847), conducted at the Alberta Children’s Hospital. This study aimed to identify neuroimaging biomarkers of recovery in children following mTBI. There was a two-stage consent process. Participants with a medically diagnosed mTBI (defined using the American Academy of Neurology criteria) were recruited through the Emergency Department and gave consent to follow-up at 4 weeks postinjury. At this time, a clinical interview and examination was performed by a physician experienced in concussion assessment where proper consent was obtained. We applied the following exclusion criteria: (1) a previous concussion within 3-months; (2) a more moderate to severe head injury (e.g. Glasgow Coma Scale (GCS) less than 13); (3) significant medical or psychiatric history; (4) medications that likely affect participation in neuroimaging and/or sleep; and (5) inability to complete questionnaires and/or neuropsychological evaluation. The final cohort thus included 110 children (8.5–17.9 years old) recovering from mTBI. None of the participants were taking any study medications at the time of imaging. For a comparative assessment (see Analysis approach section), we also collected data in an age-matched control group (n = 20, mean age = 14.44 years (standard deviation = 3.0 years), with 9 males and 11 females, see Table 1 in Iyer et al. (2019) for further demographic details). Ethics approval was granted by the University of Calgary (Canada) and the University of Queensland (Australia). Written consent from parents and child assent was obtained for all participants.

**Clinical and behavioral measures**

We considered the following clinical and behavioral measures: (1) Post–Concussion Symptom Inventory (PCSI, parent proxy and child reports), which are standardized 26-item questionnaires used to summate key domains of postconcussive symptoms and behaviors. Specifically,
was used to assess the possibility of predicting outcome
complementary support vector machine (SVM) regression
trained to predict these dichotomous outcomes, while a

Subject demographics and neuroimaging summary characteristics.

<table>
<thead>
<tr>
<th>Sample characteristics</th>
<th>Statistics</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>mTBI (n = 99)</td>
</tr>
<tr>
<td>Age</td>
<td>14.5 (2.4)</td>
</tr>
<tr>
<td>Gender</td>
<td>—</td>
</tr>
<tr>
<td>Male/female</td>
<td>48/51</td>
</tr>
<tr>
<td>&lt;14.5 years</td>
<td>28/18</td>
</tr>
<tr>
<td>≥14.5 years</td>
<td>20/33</td>
</tr>
<tr>
<td>Symptoms</td>
<td>—</td>
</tr>
<tr>
<td>Sleep</td>
<td>1.80 (0.9)</td>
</tr>
<tr>
<td>Total PCSI</td>
<td>18.2 (0–75)</td>
</tr>
<tr>
<td>Behavioral measures</td>
<td>—</td>
</tr>
<tr>
<td>CNS VS Neurocognition index (NCI)</td>
<td>98.20 (11.4)</td>
</tr>
<tr>
<td>Anatomical measures</td>
<td>—</td>
</tr>
<tr>
<td>Estimated total intracranial volume (TICV, cm³)</td>
<td>1470.60 (132)</td>
</tr>
<tr>
<td>Gray matter (GM, cm³)</td>
<td>833.60 (83)</td>
</tr>
<tr>
<td>White matter (WM, cm³)</td>
<td>432 (54.6)</td>
</tr>
<tr>
<td>Functional measures</td>
<td>—</td>
</tr>
<tr>
<td>Frame-wise displacement, FD Power (mm)</td>
<td>0.15 (0.04)</td>
</tr>
</tbody>
</table>

A summary of sample size, age and gender distribution for the two clinical groups at 1-month postinjury (recovered and symptomatic PPCS), with median and standard deviation (SD) where appropriate. Gender and age distribution in each mTBI group, split by median age, is also provided. Regarding symptoms, the median symptom scale and PCSI (Likert-scale) for the whole cohort are shown. CNS VS: Computerized Neurocognitive Software Vital Signs, with the Neurocognition index as a standard score. Anatomical measures of total intracranial volume, gray matter (GM) and white matter (WM) reported across groups. Quality control measurements (FD) from rs-fMRI were also examined. The Chi-Square test was used to assess putative group difference in Age whereas the Wilcoxon rank sum test was used to assess differences in symptoms between asymptomatic and symptomatic groups. “—” indicates nonsignificant values, *pFDR < 0.05, **pFDR < 0.001, controlling for false discovery rate (FDR).

total sleep scores were estimated using a subdomain of the PCSI that assessed a subject’s level of drowsiness, less/more sleep than usual, trouble falling asleep, and increased fatigue; and (2) a computerized cognitive assessment battery, CNS Vital Signs. Here the Neurocognition Index (NCI) was used as an overall performance index of cognitive and attentional processes.27 Subject demographics are summarized in Table 1.

Recovery was assessed at 4–6 weeks (time point, TP1) and 8–10 weeks postinjury (TP2) using overall change on the PCSI and a clinical interview conducted by a physician. Based on this interview and PCSI measurement, participants were then categorized into “Symptomatic” and “Recovered” groups. Children were considered as “Symptomatic” if there was a 10-point or greater increase in total PCSI score compared to preinjury level. Participants were considered as “Recovered” when their PCSI score returned to preinjury level (see Fig. S1, for study CONSORT diagram).28 We here refer to asymptomatic status as to those children that recovered following mTBI. Following data quality control (see below), we included 69 children in the “Symptomatic” group and 31 children in the “Recovered” group. A machine learning classifier was trained to predict these dichotomous outcomes, while a complementary support vector machine (SVM) regression was used to assess the possibility of predicting outcome along a continuum (i.e. change in PCSI and NCI scores from 4 to 8–10 weeks postinjury). Processing and analyses of neuroimaging (provided below) was conducted on imaging data collected at TP1.

Neuroimaging

Images were acquired in the oblique axial plane using a 3.0 T GE scanner (Healthcare Discovery MR, 750w, 32-channel head coil). A structural scan (T1) was acquired using the following parameters: 0.8 mm slice thickness, flip angle = 10°, inversion time = 600 ms, FOV = 240 mm. Visual inspection of each T1 scan by a radiologist and/or neuroradiologist excluded obvious signs of contusion or bleeding in gray and white matter. Resting-state functional magnetic resonance imaging (rs-fMRI) was recorded for 5 minutes and 10 seconds via gradient echo planar imaging (scanning parameters included: EPI, TE = 30 ms, TR = 2000 ms, flip angle = 90, FOV = 230 mm, 64 × 64 matrix, slice thickness = 3.6 mm). The rs-fMRI acquisition time achieved a trade-off between the need for sufficient data and the necessity of minimizing movements and burden for unwell children.29 As per described in our recent report,23 head motion (Friston 24), linear trends, and signals from cerebrospinal fluid and white matter were removed using a general linear
model framework. Data were bandpass filtered between 0.01 and 0.1 Hz. We excluded 10 subjects with less than 95% of data remaining after the removal of contaminated volumes (FD > 0.4 mm, including one preceding and two following volumes were censored). We also excluded one child due to incomplete clinical measures. Thus, we excluded 11 children from a cohort of 110, yielding a final sample size of 99 children (48 males and 51 females, no significant differences in gender). A summary on sample size characteristics and preliminary neuroimaging preprocessing details are provided in Table 1. As we used the same control cohort in our previous study, we did not exclude any healthy controls. Table 1 provides an additional summary of neuroimaging characteristics in our final sample size.

### Analysis approach

Please amend to: We started by testing for a whole-brain association between structural (voxel-based morphometry, VBM) and functional (regional homogeneity, ReHo; functional connectivity, FC) brain indices with total sleep scores. An overview of these initial analyses is shown in Figure 1. Machine learning approaches (support vector machine, SVM) were subsequently implemented in an independent group to assess the ability of the selected brain indices to predict recovery outcomes that were unbiased by total sleep scores collected at TP1. Critically, for our SVM approaches we excluded total sleep scores from the PCSI inventory. Indices of total sleep score, however, were not significantly correlated with changes in cognitive (NCI) scores adopted for the regression SVM ($r = 0.13, P = 0.18$).

We adopted an SVM approach to predict recovery outcomes, as this method offers several advantages over simpler methods including linear regression: (1) robustness to overfitting, particularly with high dimensional spaces, (2) the use of Gaussian kernel scaling functions, which enables optimized thresholding of nonlinear decision boundaries. This is critical for later assignment of decision labels (e.g. classifying Recovered vs. Symptomatic), and (3) appropriate margin of error in edge cases for labeling decisions (i.e. if a subject has opposing effects of one brain measure over other measures, SVM will maximize its decision margin to minimize error). Using a train-test approach, we trained our brain indices within an SVM classifier on 85% of subjects and tested the classifier’s accuracy on the remaining held-out individuals. This splitting of groups is in line with standard approaches in SVM classification, where data are “trained” on a larger group (usually between 80% and 90% of the sample) and “tested” on a smaller number of individuals. The selection of brain indices and training of the classifiers were therefore based on a sample of 85 individuals, with 14 individuals used as a naïve test sample to establish classification accuracy. In other words, the 14 individuals used to establish the trained classifier’s accuracy were not used to examine associations with our clinical variables, nor guide the selection of brain indices or classifier training.

Classification was repeated 10 times (10-fold cross-validation), with different subgroups comprising a similar ratio of symptomatic and recovered individuals ($n = 85$ and $n = 14$; classifier results summarized in the Supplementary Materials). All preprocessing and analyses were performed within MATLAB (MathWorks, Natick, Massachusetts), using SPM12, DPARSFA toolboxes (V4.4) and custom scripts. Rendered 3D brain visualizations for our figures were generated using MRICroGL.

### VBM

A T1-weighted scan and VBM were used to define morphological markers, Figure 1A. This was achieved by spatially normalizing all T1 images to the same stereotactic space and segmenting brain volumes into gray matter (GM), cerebrospinal fluid (CSF) and white matter (WM) images. A critical consideration for VBM in our study was the broad neurodevelopmental range (children aged between 8.5 and 18 years). We optimized VBM processing to account for variations in gray matter volume, brain size, and cortical thickness that are commonly associated with middle childhood and adolescence. Prior to segmentation of brain volumes, we first created customized tissue probability maps (TPMs), to appropriately reflect age and gender differences. Specifically, our TPM maps were created by using pediatric neuroimaging templates available from the NIH MRI study of Normal Brain Development. The age range (asymmetric templates, 4.5–18.5 years) and features present within these template images were chosen to appropriately reflect natural variations of cortical thickness and gray matter over the course of childhood. To generate these TPMs we used the Template-O-Matic (TOM8) toolbox, so that age groups could be segmented with appropriate TPMs—i.e. a TPM could be allocated for each year between 8 and 18 years.

Using age-appropriate TPMs, we next segmented brain volumes into gray matter, CSF and WM via the SPM12 toolbox using default segmentation parameters and affine regularization with an average sized template to maximize spatial accuracy within age groups. A data quality check was performed on all brain volumes. Next, diffeomorphic anatomical registration was performed to iteratively model brain shape differences arising from GM and WM images and optimize accuracy of intersubject alignment whilst correcting for volume changes. All images were...
then spatially normalized to derive smoothed (modu-
lated), weighted, Jacobian scaled gray matter images local-
ized to the Montreal Neurological Institute (MNI) space. We
employed an isotropic Gaussian kernel at 12 mm
full-width at half maximum (FWHM) for smoothing,
whilst specifying a voxel size of 3 mm for spatially nor-
malized images. As per our segmentation approach, we
performed image normalization within specific age groups
(e.g. 8–9 years, 9–10 years etc.) to limit false positive
errors that may be generated in larger group sizes.39
Smoothed images were checked across age groups for
consistency (N, median \(N=11\) subjects, across 10 age
groups). Prior to statistical analysis, all gray matter vol-
umes were then processed with an absolute threshold
mask of 0.2, to censor the possible influence of edge-ef-
ects present in other tissues (e.g. white matter).
A general linear model (GLM) was used to examine the
relationship between gray matter volume and total sleep
score. Results were corrected for family-wise error at clus-
ter-level (\(p_{FWE} < 0.05\), search threshold \(p_{uncorr} < 0.001,\)
cluster-level threshold \(k_E > 500\)). Statistical tests were
also reperformed on smoothed images processed at a
smaller kernel size (FWHM, 9 mm) with the same
family-wise error correction parameters to validate our
findings.39

ReHo (intraregional connectivity)

Processing of rs-fMRI data is specified in our recent
report.23 Importantly, for ReHo analysis described herein
unsmoothed data were used. Briefly, ReHo analysis pro-
vides a validated metric of local intra–regional functional
connectivity40 calculated by assessing the temporal syn-
chronization (or also known as Kendall coefficient of con-
dordance, KCC) of the blood-oxygen-level-dependent
(BOLD) signal between a small \(n = 27\)41 set of neighboring voxels \((n = 27)\). In this study, whole-brain ReHo
estimates provide a multimodal measure that links local
VBM estimates with between region functional connectiv-
ity. Whole-brain ReHo estimates were used to assess
brain-symptoms relationships \((p_{FWE} < 0.05\) at cluster-
level, search threshold \(p_{uncorr} < 0.005)\).

FC (interregional connectivity)

In our prior study,23 we adopted a whole-brain parcella-
tion42 to examine changes in functional connectivity pre-
sent within canonical resting-state networks in children
with mTBI. In this study, guided by our VBM and ReHo
results, we utilize a seed-based approach for measuring
interregional FC (Fig. 1C). We placed two spherical seeds

Figure 1. Processing pipeline for structural and functional data. (A) Voxel-based morphometry (VBM) was performed using structural T1 scans
from each child. Customized age-matched templates were used to derive GM, CSF and WM segmentations, before images were smoothed,
weighted and normalized to MNI space. These normalized images were then used to assess the association between gray matter volume and a
clinical construct assessing sleep disturbances. (B) Regional homogeneity (ReHo) assessment of functional imaging data (blood-oxygen-level-
dependent, BOLD, data from echo planar imaging, EPI) was used to derive intra–regional connectivity within a set of neighboring voxels \((n = 27)\). (C) Inter–regional functional connectivity was calculated using Pearson’s correlation between resting-state BOLD signals in two brain regions of
interest (posterior cingulate cortex (PCC) and medial prefrontal cortex (mPFC)).
within the brain right-hemisphere: (1) the posterior cere- 
gulate cortex (PCC, MNI coordinates: $x = 15$ mm, 
$y = -57$ mm, $z = 24$ mm; 8 mm radius sphere) and (ii) 
the medial prefrontal cortex (mPFC, MNI coordinates: 
$x = 11$ mm, $y = 58$ mm, $z = 6$ mm, 8 mm radius 
sphere). All subjects’ Pearson correlation coefficients ($r$) 
were calculated using BOLD time series from the PCC 
and mPFC. The resulting $r$-values were subsequently 
Fisher-Z transformed. Individual FC values were subse-
sequently used to perform correlations with our clinical 
variables.

**Machine learning**

We next sought to investigate the utility of the structural 
and functional brain indices to predict follow-up me-
asures of cognition and recovery. SVM regression$^{43}$ 
was utilized to determine the ability of the selected brain 
indices (VBM, ReHo, and FC) to predict change in PCSI 
and NCI scores.

An SVM classifier$^{43}$ was also performed on 85 children 
to determine if brain indices of interest could be used to 
discriminate recovery from dichotomous outcomes of 
PPCS (recovered vs. symptomatic). SVM classification 
was used to predict dichotomous clinical outcomes via a 
hold-out cross-validation ($n = 14$ used as test sample). 
The SVM accuracy was further tested using 10-fold cross-
validation in nine additional data groupings (i.e. each 
additional group comprised 85 individuals as training 
samples and 14 individuals as naive test samples). The 
average performance of both SVM regression and SVM 
classifier results are presented in the Results section.

**Results**

**DMN structure and function in children with 
PPCS is related with sleep disturbances**

We found a significant negative association between PCC 
and mPFC gray matter volumes and sleep scores: Gray 
matter volume in these brain regions was lower in chil-
dren with higher sleep disturbances ($n = 85$, 
$r = -0.30 \pm 0.008$ standard error (SE) across 10-folds, 
$p_{FWE} = 0.002$ for PCC, $r = -0.38 \pm 0.013$ SE, 
$p_{FWE} = 0.0001$ for mPFC; Fig. 2A). Negative associations 
between gray matter volumes and total sleep scores were 
also confirmed for spatially smoothed at a kernel size of 
9 mm (across 10-folds, $r = -0.34 \pm 0.012$ SE for PCC

Figure 2. Whole-brain association of structure and function with symptoms. (A) Gray matter volume changes, detected using voxel-based 
morphometry (VBM), negatively correlated with sleep scores such that decreased gray matter volume in the right PCC and the right mPFC was 
linked to increased sleep disturbances and fatigue (cluster-level $p_{FWE} < 0.002$ and $p_{FWE} < 0.0001$, respectively; high threshold of $p_{uncorr} < 0.001$). 
(B) Reduced within-region functional connectivity (ReHo) in the right PCC also negatively correlated with sleep problems ($p_{FWE} = 0.02$ at cluster-
level, high threshold of $p_{uncorr} < 0.005$). (C) Functional connectivity (FC) between the right PCC and mPFC indicates a negative correlation: 
Children with lower across-region FC showed increased sleep problems. Results in this figure are from the first data grouping (fold) 1. All 
correlations were adjusted for age, gender and brain volumes.
ReHo in the PCC was negatively associated with sleep problems \( (r = -0.25 \pm 0.036 \, \text{SE}, \, P_{FWE} = 0.02) \); Fig. 2B). In line with the aforementioned gray matter volume associations, decreased ReHo in the PCC was related to increased sleep problems.

In agreement with the associations between local brain indices and sleep-related scores, FC between the PCC and mPFC also negatively correlated with sleep scores \( (r = -0.28 \pm 0.02 \, \text{SE}, \, P = 0.006) \). Across our measures, we only observed an effect of gender for the correlation between measures of FC between the PCC and mPFC and sleep scores (female \( r = -0.28 \pm 0.02 \, \text{SE}, \, P = 0.006 \), male children \( r = -0.08 \pm 0.03 \, \text{SE}, \, P = 0.61 \)).

Following this observation, we investigated possible differences in sleep symptom severity between genders. The results showed higher sleep disturbances in females compared to males (unpaired \( t \)-test, \( t_{X7} = 3.20, \, P = 0.002 \). However, values of FC were similar between genders (unpaired \( t \)-test, \( t_{X7} = 1.55, \, P = 0.17 \)). See Table S1 for a summary of associations and the relative contribution of age, gender and brain volumes.

Additionally, the comparison between brain indices of mTBI children with our age-matched controls revealed group differences in gray matter and ReHo measures, but not in FC (one-way ANOVA we report significant differences for GM in PCC: \( P_{FDR} = 1.1 \times 10^{-12} \), GM in mPFC: \( P_{FDR} = 1.2 \times 10^{-9} \); ReHo in PCC: \( P_{FDR} = 5.3 \times 10^{-7} \); FC: \( P_{FDR} = 0.25 \), see Fig. S2).

**Brain signatures of sleep carry prognostic information**

SVM regression included the four brain indices collected at 1-month postinjury: gray matter volume (eigenvariate) extracted from the PCC and mPFC, ReHo estimates from the PCC, and PCC–mPFC FC values. Age at 1-month postinjury was also included. Changes in PCSI scores (8–10 weeks follow-up scores minus 4 weeks postinjury scores; total sleep scores excluded) predicted by these five indices were highly correlated with real PCSI scores (10-folds average: \( r = 0.52, \, P = 0.0009, \, 12.28\% \text{ standard error} \); Fig. 3A). Moreover, SVM regression using the same brain indices was also able to predict changes in cognitive (NCI) scores (10-folds average: \( r = 0.43, \, P = 0.0002, \, 4.40\% \text{ standard error} \)).

The SVM classifier, adopting the abovementioned four brain features plus age, was able to accurately discriminate children who recovered from concussion to those who remained clinically symptomatic (10-folds average: 86% accuracy, 69% specificity and 94% sensitivity; Fig. 3B). The accuracy range for the classifier to discriminate symptomatic from recovered children was between 78% and 92% (10-folds; Fig. 3B). The prediction accuracy on our hold-out test samples \( (n = 14) \) also maintained high accuracy (10-folds average: 79% accuracy, 75% specificity, and 82% sensitivity). The range of accuracy in the test samples was between 65% and 95% (10-folds). See Tables S2 and S3 for a summary of classification results.

**Figure 3.** Classification and outcome assessment. (A) SVM regression was performed using four brain features linked to behavioral sleep scores at 1-month postinjury: PCC and mPFC gray matter volumes, PCC values of local functional connectivity (ReHo), and resting-state functional connectivity between PCC and mPFC. These brain features, plus the age of the participant, were able to significantly predict changes in post–concussion symptoms between 1-month postinjury and 8–10 weeks follow-up along a continuum. PCSI = Post–Concussion Symptom Inventory. (B) SVM classification results using the four brain features of interest and age of the child to classify recovery versus symptomatic PPCS at follow-up (8–10 weeks mark). The receiver operating characteristic curve (ROC) shows an area under the curve (AUC) with high specificity and sensitivity. The confidence interval (dashed lines) indicates the most and least accurate classification across the 10-folds. (C) Accuracy of age and each brain feature (GM, ReHo and FC) – tested independently – in predicting recovery outcome. Error bars indicate averages over 10-folds (±standard deviation).
Discussion

This study supports the notion that structural and functional brain changes in the DMN underpin core manifestations of pediatric PPCS following mTBI. Our findings also extend upon previous work by showing that indices of symptom-related changes in gray matter and functional connectivity within and between two key regions of the DMN — PCC and mPFC — carry significant prognostic information.

Prior studies in both children and adults have suggested that the DMN plays a crucial role in sleep processes, which in turn helps maintain cognitive performance. Accordingly, it has been shown that TBI causes structural and functional changes within the DMN that have been linked to cognitive deficits, including slower rates of information processing and decision-making. Though the neural underpinning of poor sleep and excessive fatigue following mTBI has not been elucidated by prior work, recent studies in adolescents have pointed towards weaker DMN functional connectivity.

Our study directly supports and extends these emerging evidences by showing that converging decreases in gray matter volume, as well as intra- and inter-regional functional connectivity, in the PCC and mPFC in children with mTBI-induced PPCS are linked to increases in sleep disturbances and fatigue. Negative associations of gray matter and intraregional connectivity with sleep-related problems occurred within the context of higher brain index values compared to matched healthy controls (Fig. S2). These findings provide support to the notion of a neurotrauma-related compensatory response within DMN regions. Although gray matter and intra–regional connectivity values did not indicate gender-specific associations, the negative association between interregional FC and sleep disturbances was driven by females. This association, along with the fact that females reported higher levels of sleep-related symptoms compared to males, is in line with the suggestion that concussed females are more sensitive to sleep difficulties and have protracted recovery patterns.

Cognitive functions, including reasoning and working memory, have been associated with higher DMN functional connectivity at rest. Accordingly, poorer cognitive functions following TBI have been linked to reduced DMN structural and resting-state functional connectivity. In our previous exploratory analysis conducted on the same cohort, we were unable to detect a significant change in global DMN resting-state functional connectivity in children following mTBI compared to matched controls. We did, however, find an association between variations in DMN global connectivity and cognitive symptoms. The current reanalysis of the data offers an important insight into our previous results, indicating that key symptoms of PPCS following mTBI are linked to circumscribed changes to the structure and function of two key DMN nodes. Our linear classifiers build upon this assertion, showing that a combination of PCC and mPFC structural and functional features can significantly and accurately predict cognitive outcomes. DMN functional connectivity in children has shown to normalize following improvements in cognitive functions and sleep quality. Our findings are consistent with these previous reports and highlight that a normalization of functional interactions between, as well as within, the PCC and the mPFC is most likely responsible for the observed behavioral recovery.

Recovery from PPCS in children is complex to predict. Monitoring of symptoms and clinical assessments using scales like the PCSI have shown some sensitivity in predicting behavioral outcomes. However, the predictive value of psychometric and behavioral measurements in tracking recovery trajectory alone is insufficient, limiting the ability to develop and implement outcome-oriented personalized therapeutic interventions. Our findings motivate the use of neuroimaging indices to supplement existing clinical evaluations of symptoms and behaviors during recovery from PPCS. Specifically, the anatomical and functional assessment PCC and mPFC provide useful insights on the likelihood of recovery of a developing injured brain. In fact, our findings suggest that convergent anatomical and functional changes within and between these two DMN regions offer valuable prognostic information on the temporal evolution of core PPCS. These results echo prior neuroimaging studies showing that improvement in adult recovery following TBI is linked with higher activity within the DMN.

Predicting the recovery trajectory of a child with PPCS is encumbered by the high inter–individual variability in symptoms, age, gender, and premorbid characteristics. Our findings showed that the inter–individual variability of PPCS recovery in children is linked to variations in brain indices that are indexed by relatively simple clinical scores of sleep disturbance and fatigue. In this context we note that, while appealing, a dichotomous classification of outcomes (recovered vs. symptomatic) may misrepresent the actual progression of PPCS following mTBI. In effect, our machine learning results highlight that changes in PPCS including cognitive functions can be predicted by an SVM regression approach. That is, the collection and assessment of anatomical and functional brain indices alongside standard neuropsychological assessments at different timepoints postinjury offer potential for prognosis along a natural continuum of recovery. More broadly, our results are in agreement with the suggestion that recovery from mTBI maps onto a dynamic and “multifaceted construct”.

In sum, our study complements prior work in supporting the key role of sleep and fatigue to the clinical picture of PPCS. The significant association between variations in such key symptoms and structure-function in PCC and mPFC directly support the notion that the DMN underpins core manifestations of mTBI-induced PPCS. The prognostic value of these brain indices further encourages this hypothesis and provides a strong rationale for translational efforts aiming to facilitate the use of neuroimaging in clinical settings. Future validation studies on independent datasets are required to test and improve the predictive utility of the brain indices identified in this study towards overall recovery outcomes. The provision of additional information gathered from neuroimaging may indeed enable the targeted allocation of limited therapeutic resources to children with a greater likelihood of poor outcome.

Acknowledgments

K.K.I acknowledges funding support from the Brain Foundation. K.M.B. acknowledges funding support from the Motor Accident Insurance Commission (MAIC, Queensland, Australia) and the Canadian Institutes of Health Research (grant number: 293375). L.C. is supported by the Australian National Health Medical Research Council (L.C. 1099082 and 1138711). We thank Nathan Stevenson for his advice on implementation of support vector machine classification with our dataset.

Conflict of Interest

The authors have no competing interest or conflict of interest to declare.

Author Contributions

K.K.I. performed all aspects of preprocessing, analysis, interpretation of results, writing of the manuscript and evaluation of findings. A.Z. assisted with analysis, statistical inferences, writing of the manuscript and evaluation of findings. K.M.B conceived, designed, acquired data for the study and provided clinical interpretation of the findings. L.C. performed all aspects of preprocessing, analysis, interpretation of results, writing of the manuscript and evaluation of findings.

References

22. Davey CG, Harrison BJ. The brain’s center of gravity: how the default mode network helps us to understand the self. World Psychiatry 2018;17:278.

Supporting Information
Additional supporting information may be found online in the Supporting Information section at the end of the article.

Figure S1. Study CONSORT diagram.
Table S1. General linear modeling results.
Figure S2. Brain indices in mTBI and age-matched healthy controls.
Table S2. Classification results (SVM regression).
Table S3. Classification results (Outcome classifier).