Multiparametric MRI changes persist beyond recovery in concussed adolescent hockey players

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ABSTRACT

Objective: To determine whether multiparametric MRI data can provide insight into the acute and long-lasting neuronal sequelae after a concussion in adolescent athletes.

Methods: Players were recruited from Bantam hockey leagues in which body checking is first introduced (male, age 11–14 years). Clinical measures, diffusion metrics, resting-state network and region-to-region functional connectivity patterns, and magnetic resonance spectroscopy absolute metabolite concentrations were analyzed from an independent, age-matched control group of hockey players (n = 26) and longitudinally in concussed athletes within 24 to 72 hours (n = 17) and 3 months (n = 14) after a diagnosed concussion.

Results: There were diffusion abnormalities within multiple white matter tracts, functional hyperconnectivity, and decreases in choline 3 months after concussion. Tract-specific spatial statistics revealed a large region along the superior longitudinal fasciculus with the largest decreases in diffusivity measures, which significantly correlated with clinical deficits. This region also spatially intersected with probabilistic tracts connecting cortical regions where we found acute functional connectivity changes. Hyperconnectivity patterns at 3 months after concussion were present only in players with relatively less severe clinical outcomes, higher choline concentrations, and diffusivity indicative of relatively less axonal disruption.

Conclusions: Changes persisted well after players’ clinical scores had returned to normal and they had been cleared to return to play. Ongoing white matter maturation may make adolescent athletes particularly vulnerable to brain injury, and they may require extended recovery periods. The consequences of early brain injury for ongoing brain development and risk of more serious conditions such as second impact syndrome or neural degenerative processes need to be elucidated.

Neurology® 2017;89:2157–2166

GLOSSARY

AD = axial diffusivity; CST = corticospinal tract; DMN = default mode network; DTI = diffusion tensor imaging; FA = fractional anisotropy; MD = mean diffusivity; MRS = magnetic resonance spectroscopy; ROIs = region of interest; RS = resting-state; RSN = resting-state network; SLF = superior longitudinal fasciculus.

Concussion remains a critical health problem, and the developing brain may be uniquely susceptible. In Canada, hockey-related concussions account for nearly half of all sports team-related concussions and disproportionately affect adolescents. The neuronal mechanisms that underlie concussion remain poorly understood. A complex cascade of structural, functional, and metabolic changes are secondary to rotational acceleration of the brain. MRI studies have reported neuronal vulnerabilities that persist after athletes have been cleared to return to play. Without full recovery, players may have increased susceptibility to injury and may be at risk of serious conditions such as second impact syndrome. Therefore, it is essential to understand the
immediate effects of concussion on the adolescent brain and the nature of any long-lasting neuronal changes detectable with noninvasive multiparametric imaging.

In this study, concussed adolescent male hockey players were evaluated longitudinally after concussion and compared to independent, age-matched nonconcussed players as controls. While clinical measures in this population indicate recovery and prompt return to play, given previous imaging findings, we hypothesized that there would be persistent MRI-detectable neuronal changes that reflect long-lasting damage to white matter fibers and compensatory recovery mechanisms. Given this multiparametric MRI dataset, we were able to directly relate resting-state (RS-fMRI) changes with underlying structural abnormalities through diffusion tensor imaging (DTI), brain metabolic changes through magnetic resonance spectroscopy (MRS), and acute clinical deficits.

METHODS

Players diagnosed with a mild traumatic brain injury or concussion (age 13.3 ± 0.6 years) were assessed within 24 to 72 hours of their injury (n = 17) and again 3 months later (n = 14). An independent group of healthy, age-matched, nonconcussed male hockey players were assessed and acted as our controls (n = 26, age 13.0 ± 1 years). Portions of this study design have been described previously.14 Further clinical methods are described in appendix e-1 at Neurology.org. The numbers of included participants for each portion of the study are summarized in table e-1. MRI acquisition parameters are outlined in appendix e-2.

Standard protocol approvals, registrations, and patient consents. Ethics approval was obtained through the University of Western Ontario's Health Sciences Research Ethics Board. All participants and parents/guardians provided informed written consent.

RS-fMRI analysis. RS-fMRI data were analyzed with FSL (FMRIB Software Library) 5.0.6 (Oxford, England) with 2 major aims: to investigate RS network (RSN) reorganization and whole-brain region-to-region connectivity patterns. Standard preprocessing steps included brain extraction, a high-pass filter (0.01 Hz), spatial smoothing with a 5-mm full-width half-maximum gaussian filter, and motion correction using linear image registration relative to the middle volume. Data with excessive motion (>1-mm maximum displacement or >0.5-mm relative mean displacement) were excluded from further analyses. Preprocessed images were denoised with single-session independent component analysis and automatic dimensionality estimation. Components that were designated as noise (related voxels outside the brain, within the corticospinal fluid, primarily high-frequency profile, or motion artifacts) were regressed from the data. Each dataset was registered to standard space with a 2-stage affine registration technique that first registered the low-resolution functional data to the brain-extracted anatomic image and then to a 2-mm isotropic Montreal Neurologic Institute T1-weighted reference image using 12 df.

Multisession temporal concatenation–independent component analysis was used to create average RSN maps followed by dual regression to generate session-specific networks.15 In particular, the visual occipital pole, default mode (DMN), sensorimotor, executive control, and cerebellar networks were investigated. The FSL randomize permutation-testing tool was implemented to assess group differences according to a general linear model with threshold-free cluster enhancement and multiple-comparison Bonferroni-corrected p values (p < 0.001). In addition, a regional connectivity analysis was performed on the preprocessed data with the CONN toolbox because of the statistical advantages as it complements the network analysis and expands to look at whole-brain regional connectivity. Participant-specific CSF and white matter masks were used as regressors. Data were parcellated into 136 functionally relevant regions with the Harvard-Oxford cortical atlas and cerebellar regions with the Automated Anatomical Labeling atlas. Whole-brain region-to-region connectivity differences were assessed with false discovery rate correction.

Diffusion tensor and tractography analysis. Raw diffusion datasets were corrected for eddy current distortions and relative head motion with the eddy command line tool. The non-diffusion-weighted (b = 0) volume was used to create a modestly extracted brain mask. A diffusion tensor was fit voxel-wise with the use of specific inputs of the gradient directions and b values (b = 1,000 s/mm²) directly from the scanner. Several diffusion metric spatial maps were created, including fractional anisotropy (FA) and mean (MD), axial (AD), and radial (RD) diffusivity. With the use of the John Hopkins University high-probability white matter tractography atlas,13 these DTI metrics were extracted and analyzed within the superior longitudinal fasciculus (SLF), cingulum, forceps major and minor of the corpus callosum, uncinate fasciculus, corticospinal tract (CST), anterior thalamic radiation, and inferior fronto-occipital and longitudinal fasciculus.

Once tracts with significant changes were identified with a multivariate analysis of variance, an iterative randomization tool within FSL was used to spatially quantify where the greatest changes occurred along those tracts while regressing for age. DTI metrics within the regions with the greatest changes were extracted and analyzed. These regions are referred to as DTImax and are described in Results. Probabilistic tractography with iterative Monte Carlo simulations of principal diffusion tensor vectors was used to quantify a probabilistic streamline between regions of interest (ROIs). Probabilistic structural connections were investigated between relevant parcelized RS-fMRI regions to spatially relate structural and functional changes. To quantify these results, 2 identical spherical ROIs (5-mm radius) were placed before and after DTImax (that overlapped along the probabilistic tract connecting acute RS-fMRI ROIs that we identified, figure 1A). The probability distribution was displayed per participant and quantified the probability of a structural connection between the seed and waypoint masks (normalized by the total number of tracts modeled).

MRS analysis. Spectra were postprocessed with in-house software to measure absolute N-acetylaspartate, choline, creatine, glutamate, glutamine, and myo-inositol as previously described. Briefly, analysis software created in our laboratory in IDL (version 5.4, Research Systems Inc, Boulder, CO) was used to correct spectral line shapes by combined QUALITY deconvolution and eddy current correction. Spectra were fitted in the time domain with the use of a Levenburg-Marquardt minimization routine with prior knowledge of metabolic line shapes. Prior knowledge was acquired from in vitro spectra obtained from aqueous solutions of metabolites before the study. We report absolute metabolite levels using unsuppressed
water from the ROI as an internal standard as previously described in detail.\textsuperscript{14,16} In addition, absolute concentrations included a correction for tissue partial volume and T1 and T2 relaxation–related signal loss.

**Statistical analysis.** Statistical analysis was performed with a multivariate analysis of variance with Tukey post hoc testing using GraphPad Prism (version 6.0). A hypothesis-driven 2-tailed Pearson correlation analysis was explored between data that had a significant effect for group ($p$, 0.05 after false discovery rate correction).

**RESULTS** Clinical results are summarized in appendix e-3 and figure e-1.

**RS-fMRI results.** We found significant increases in connectivity at 3 months after concussion in several RSNs. In particular, there was significantly increased connectivity at 3 months after concussion compared to controls with the occipital pole visual network and cerebellar network (figure 2, B and C). Compared to 24 to 72 hours after concussion, there were significant increases in connectivity by 3 months with the somatosensory network (figure 2D) and cerebellar network.

There was increased DMN connectivity at 3 months compared to controls that did not survive statistical Bonferroni correction ($p < 0.01$, figure 2A), and no significant differences were found within the executive control network or between controls and the 24- to 72-hour postconcussion group in any network.

There were a few subtle changes in region-to-region connectivity within 24 to 72 hours after concussion compared to controls (figure 1B); however, some were striking with a large effect size (>0.8) and prompted further analysis to directly relate with our DTI findings (figure 1C). Specifically, there were significant anticorrelations between the anterior supramarginal gyrus and areas that mimic the DMN, including the angular gyrus and frontal pole ($p$, 0.05, including portions of the medial prefrontal cortex, figure 1C). This was not found in our network analysis, possibly as a result of statistical power (voxel-level correction) or the dominance of the frontal pole. There was an increase in regional connectivity immediately after concussion compared to controls.
between the cerebellum and parahippocampal gyrus. The significant whole-brain region-to-region connectivity changes are shown in figure 3, and similar to the RSN results, there were increases in connectivity at 3 months compared to both the control and 24- to 72-hour postconcussion groups between regions that complemented our network-level connectivity changes.

Diffusion results. There was a significant main effect for group differences in the CST, cingulum, and SLF (F > 4.18, p < 0.05, figure 4). Along the CST, the greatest diffusion changes were located inferiorly near the brainstem (p < 0.1), and there were small, more scattered maximums along the cingulum (p > 0.1). A large region along the right SLF shown in figure 1 had significant MD, RD, and AD changes at 3 months after concussion (p < 0.01), and a mask was created over this localized hotspot of compromised neuronal integrity, DTI_max. Within DTI_max, the MD and RD values were significantly decreased at both 24 to 72 hours and 3 months after concussion compared to controls (F > 10.0, p < 0.01); furthermore, there were decreases in AD at both time points (F = 6.67, p = 0.060 and p = 0.003, respectively) with increases in FA by 3 months (F = 3.91, p = 0.025). Using the anterior supramarginal gyrus as a seed and the frontal pole as a waypoint, we confirmed that probabilistic tracts structurally connected regions where we found acute RS-fMRI differences (figure 1). The probability of a structural connection did not significantly decrease after concussion.

Spectroscopy results. The spectroscopy voxel was placed in the prefrontal region with the following mean ± SD tissue content: gray matter 19% ± 7%, white matter 78% ± 8%, and CSF 3% ± 2%. The tissue composition of the voxel did not significantly change across participants or sessions. No changes in N-acetylaspartate were found between any of the groups. Choline was significantly reduced at 3 months relative to controls (F = 4.2, p = 0.02, figure 4), with a 10% decrease in the mean observed (p = 0.035). A reduction in glutamine levels was observed but was not significantly different from controls (F = 2.3, p = 0.09). Correlation results are summarized in detail in table e-2, and examples are shown in figure 5.
DISCUSSION In this study of adolescent hockey players from Bantam leagues in which body checking is first introduced, we found acute and longitudinal multiparametric MRI changes after concussion compared to controls. As expected, clinical composite scores elevated immediately after concussion, and within weeks of injury, they returned to control levels and players were cleared to return to play. However, we found persistent RS-fMRI connectivity changes, diffusion-related white matter abnormalities, and MRS metabolite decreases in the prefrontal white matter at 3 months after concussion.

The adolescent brain is still not fully developed during this age range (11–14 years), and as it continues to grow and mature, it may be more vulnerable to brain dysfunction and elongated periods of recovery after an acceleration-related injury. It has been shown that this heightened vulnerability may be due to both biomechanical (neck strength) and neurobiological (incomplete white matter myelination) characteristics. Axonal excitability and electrophysiologic recordings in a rat fluid percussion brain injury model have been used to assess differences in axonal vulnerability between myelinated and unmyelinated fibers during injury response and functional recovery. In this animal model, myelinated axons recovered within 7 days and were relatively protected compared to unmyelinated fibers, which were more severely injured and took longer to recover. It is critical to understand how the adolescent human brain reacts and recovers from concussion. A recent review of concussion-related MRI findings demonstrated the advantages of relating multiple MRI modalities, and here we aim to do just that to better understand the meaning of these changes.

We found diffuse changes in DTI metrics along 3 major white matter tracts. However, the tract-specific spatial changes were localized to a single, central region along the SLF (DTI $\text{max}$) where we found decreased AD, MD, and RD at both times after concussion. In vivo animal studies have been able to directly relate histology with MRI diffusion, specifically finding a direct...
relationship between AD and axonal injury through postmortem neurofilament immunostaining and confirmation of the presence of β-amyloid precursor protein.21 Diffuse axonal pathology has been observed in all traumatic brain injury severities, in which transport is disrupted and axonal bulb formations occur.22 The decreases in AD observed after concussion may reflect this axonal pathology because they relate directly to immediate symptoms. The AD within DTI_{max} at 24 to 72 hours after concussion was significantly correlated with total symptom scores and the number of errors during balance testing, indicating that a more severe injury is associated with more severe axonal damage. This remained true at 3 months after concussion when the sustained decrease in AD within DTI_{max} was associated with a higher symptom severity score.

The most common injury mechanism in our cohort was falling, affecting the back of the head. Investigations of the biomechanical deformations and shearing forces after even minimal acceleration of the human head have reported deformation of the human brain. Rapid deceleration directed to the back
of the head results in anterior shortening and posterior elongation, as well as deformations in the axial plane. These results correspond to regions where we found diffusion changes indicative of structural damage to the long tracts running anterior to posterior. This acceleration injury and the consequent strain and shearing forces between brain tissues may damage axons themselves. The decrease in AD was concurrent with decreases in RD and MD and increases in FA within the same region. This could reflect different pathologies such as axonal disruption, cytotoxic edema, axonal swelling, or changes in myelin at the site of injury. These findings are similar to some previous DTI studies but not all. Cohorts of different ages, injury severity, and acquisition timing all may play a role in the direction of these changing.

Figure 5  Examples of correlation results

Relating data at (A and B) 24 to 72 hours postconcussion (PC), (C and D) 3 months PC, and (E and F) between 24 to 72 hours PC (x-axis) and 3 months PC (y-axis) data. Further details are provided in table e-2. The 95% confidence interval is shown using the dashed curves, and all relationships shown here have p < 0.05 after false discovery rate correction. AD = axial diffusivity; CHO = choline; DTI = diffusion tensor imaging; ImPACT = Immediate Post-Concussion Assessment and Cognitive Testing; MRS = magnetic resonance spectroscopy; SLF = superior longitudinal fasciculus.
diffusion metrics. Fluid-attenuated inversion recovery and turbo spin echo sequences were used to confirm a mild traumatic brain injury with no cerebral edema. We observed region-to-region RS-fMRI hyperconnectivity at 3 months after concussion compared to both controls and 24 to 72 hours after concussion. This was also true at the network level. Previous studies have variably found this pattern, and hyperconnectivity has been proposed to be involved in recovery and compensation for underlying white matter disruption.6,30–35

Here, we demonstrate the strength of multiparametric MRI by relating our connectivity findings directly to DTI and clinical data. Acute regional connectivity changes with the cerebellum correlated with all of the significant measures reflecting acute postural balance deficits and directly relate the role of the cerebellum in balance performance.6 The average AD in the cerebellum in balance performance at 3 months relative to controls. The average AD along the SLF and within DTI_max was correlated with changes in interregional connectivity in which, in general, more axonal injury (lower diffusivity) and a more severe injury as quantified by number and severity of symptoms or more days required to recover were associated with lower absolute values of connectivity. Only those players with a less severe injury as indicated by these acute clinical measures exhibited regional hyperconnectivity at 3 months after concussion, similar to previous work focused on the DMN in more severe and older patients with traumatic brain injury 6 months after their injury.55 This possible recovery mechanism may be compensating for the underlying damaged avenues of structural connections by recruiting regions of connectivity and enhancing both correlated and inhibitory communication. It is possible that some players have not yet reached or completed the recovery phase, and further longitudinal data are needed to understand the timing and potential long-term consequences of these mechanisms.

A 10% reduction in the MRS choline signal was observed at 3 months relative to controls. The choline signal is made up of different components, mainly phosphocholine (a precursor to membrane synthesis) and glycerophosphorylcholine (a breakdown product of the membrane).36 This change is particularly interesting because reductions in blood plasma glycerophospholipids at 24 to 72 hours were reported in a subset of these hockey players.10 Other studies have also reported reduced choline levels in the motor cortex after a month of high school football in non-concussed players.37 This was suggested to be due to a decrease in membrane turnover as a result of repetitive subconcussive impacts. The correlations with the MRS data support similar interpretations. The choline concentration at both 24 to 72 hours and 3 months after concussion was significantly correlated with RS-fMRI connectivity; in particular, a lower choline concentration was associated with a lower absolute interregional connectivity value. Given the relationship between choline concentrations and the clinical, diffusion, and connectivity measures, we hypothesize that the decrease in choline is related to reduced membrane turnover rate.36–39

There are limitations in the present study that will motivate future MRI investigations of adolescent concussion. The average changes we report here represent a heterogeneous population whose individual injury mechanisms, severities, and recovery times vary. We followed up with players 3 months after their injury; however, the players had clinically recovered and were cleared to return to play at different times. It is possible that returning to play may affect the brain and any ongoing neuroreparative mechanisms. Given the long-lasting changes we found at 3 months, future longitudinal studies should continue to follow this demographic as they develop to understand exactly how long the recovery period lasts as a function of age and the consequences, if any, later in life, including increased vulnerability to future concussions and the development of neurodegenerative diseases such as chronic traumatic encephalopathy.38,39

The persistent changes we observed in this study provide evidence of prolonged axonal disruption in a localized area along the SLF. This result was spatially related to acute RS-fMRI connectivity changes and choline metabolite decreases and correlated significantly with clinical deficits. Diffuse hyperconnectivity patterns were present 3 months after concussion, well after clearance to return to play and symptomatic recovery. This adolescent population may be particularly vulnerable to injury while axons continue to myelinate and mature, and our results suggest that they may require longer recovery periods.

AUTHOR CONTRIBUTIONS
All authors approved the manuscript and provided critical edits. K.Y.M.: manuscript preparation, analysis and interpretation of data, statistical analysis. A.S.: analysis and interpretation of data. R.B.: study concept and design. G.A.D.: overall project design. C.B.: study coordinator. A.B.: overall project design. L.F.: overall project design, patient evaluation. K.A.: patient evaluation. T.J.D.: interpretation of data. D.D.F.: overall project design. J.H.: overall project design, clinical acquisition. R.S.M.: overall project and image acquisition design, interpretation of data.

ACKNOWLEDGMENT
The authors acknowledge Sandra Shaw for her efforts in organizing patient appointments. They also thank and commend the players and their parents/guardians for their willingness to participate in this study and the coaches and MRI technicians for their excellent support.

STUDY FUNDING
Funding was provided by a grant from the Children’s Health Foundation to Dr. D.D. Fraser and a grant from the Western University Schulich School of Medicine and Dentistry (co-principal investigators: Dr. D.D. Fraser and G.A. Dekaban).
DISCLOSURE
K. Manning and A. Schrani report no disclosures relevant to the manuscript. R. Bartha is funded by CIHR Project Grant 152927; Canadian Consortium on Neurodegeneration in Aging 006358; Natural Sciences and Engineering Research Council of Canada grant 25033-2013; Ontario Institute for Cancer Research Grant 00870; Smarter Imaging Program, Ontario Brain Institute grant ONDRI2013, and an Alzheimer Foundation of London and Middlesex Premier Research Grant. G. Dekaban is funded by CIHR project grant PJT 144651, is co-principal investigator on the London Health Sciences Centre Children’s Health Foundation grant LHR F7806 R-13-116 CHRI (funded the work presented in this publication), is co-principal investigator or applicant on 3 additional CIHR grants, and receives funds from the NSE BioCanRx, C. Barreira, A. Brown, L. Fischer, and K. Asem report no disclosures relevant to the manuscript. T. Doherty receives research support from Allergan Canada. D. Fraser receives funding through the Children’s Health Foundation. J. Holmes is funded by CIHR operating grant 326065 and an Alzheimer’s Association International Research Grant. R. Menon is funded by CIHR Foundation Grant 353372; Natural Sciences and Engineering Research Council of Canada grant 261701-2012, and a Brain Canada Platform Support Grant; has sold a patent to Siemens Healthcare; and receives research support from Siemens Healthineers. Go to Neurology.org for full disclosures.

Received April 19, 2017. Accepted in final form September 6, 2017.

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