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Relating white matter microstructure in theoretically defined addiction networks to relapse in alcohol use disorder

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Theoretical models group maladaptive behaviors in addiction into neurocognitive domains such as incentive salience (IS), negative emotionality (NE), and executive functioning (EF). Alterations in these domains lead to relapse in alcohol use disorder (AUD). We examine whether microstructural measures in the white matter pathways supporting these domains are associated with relapse in AUD. Diffusion kurtosis imaging data were collected from 53 individuals with AUD during early abstinence. We used probabilistic tractography to delineate the fornix (IS), uncinate fasciculus (NE), and anterior thalamic radiation (EF) in each participant and extracted mean fractional anisotropy (FA) and kurtosis fractional anisotropy (KFA) within each tract. Binary (abstained vs. relapsed) and continuous (number of days abstinent) relapse measures were collected over a 4-month period. Across tracts, anisotropy measures were typically (i) lower in those that relapsed during the follow-up period and (ii) positively associated with the duration of sustained abstinence during the follow-up period. However, only KFA in the right fornix reached significance in our sample. The association between microstructural measures in these fiber tracts and treatment outcome in a small sample highlights the potential utility of the three-factor model of addiction and the role of white matter alterations in AUD.

Key words: alcohol use disorder; diffusion MRI; relapse; white matter.

Introduction

Alcohol use disorder (AUD) is a common psychiatric condition characterized by chronic alcohol use despite harmful consequences. In the United States, roughly 30% of adults will meet criteria for AUD at some point in their lives (Grant et al. 2015). Although residential and outpatient treatment programs are widely available, patients with AUD frequently relapse following treatment and are trapped in a cycle between treatment, abstinence, and relapse (McKay and Hiller-Sturmhöfel 2011). AUD's persistent relapse rates highlight the importance of developing new treatment strategies such as neuromodulation interventions that are designed with a greater understanding of the risk factors and neurobiological mechanisms involved in relapse.

AUD results from a culmination of psychosocial, environmental and biological factors (Deeken et al. 2020; Ossola et al. 2021). Among the biological factors are maladaptive alterations to neurocircuitry, which have been a primary focus of addiction research for the past several decades. Koob and Volkow synthesized this body of animal and human imaging research into a theoretical model which describes the neural pathophysiology of addiction (Koob and Volkow 2010; Koob and Volkow 2016). This framework outlines three neurofunctional domains impacted in addiction: incentive salience (IS; e.g. exaggerated reward response from a drug), negative emotionality (NE; e.g. withdrawal and exaggerated stress when not using), and executive functioning (EF; e.g. preoccupation with a substance; heightened impulsivity).

The neurofunctional domains identified by Koob and Volkow and alterations to the structures that underlie those domains may explain individual variation in resilience to relapse (Koob and Volkow 2010; Koob and Volkow 2016). We previously showed that reduced resting state functional connectivity measured during early abstinence within addiction networks defined by Koob and Volkow's theoretical model of addiction were associated with an increased risk of relapse and predicted days-to-relapse in AUD (Camchong et al. 2022). In the present manuscript, we leverage modern diffusion MRI techniques to examine white matter microstructure within white matter tracts connecting regions within these addiction networks. For each of the three Koob-Volkow domains, we identified a white matter tract based on prior research. Fig. 1 illustrates these three tracts in an example participant.

IS domain and the fornix

In AUD, the IS domain involves an exaggerated reward response to alcohol. This heightened reward response is thought to be



Fig. 1. Tractography results rendered as 3D surfaces overlaid on an FA map. The fornix (yellow), uncinate fasciculus (blue), and anterior thalamic radiations (green) correspond to the IS, NE, and EF domains, respectively.

modulated by excess dopaminergic activity in the mesolimbic reward pathway (Vollstädt-Klein et al. 2010; Sanchez-Roige et al. 2014), including the nucleus accumbens. Supporting connectivity between the nucleus accumbens and several other limbic structures (e.g. hippocampus, thalamus, and hypothalamus), the fornix is a primary white matter pathway mediating reward and memory circuitry (Brown and Winocur 1973; Shin et al. 2019). Decreased FA in the fornix has been reported in alcoholics relative to controls (Pfefferbaum et al. 2009) and fornix FA has been shown to be negatively correlated with alcohol cue reactivity in heavy drinkers (Monnig et al. 2013).

NE domain and the uncinate fasciculus

The NE domain describes how alcohol-seeking behavior in AUD is characterized by negative affect such as dysphoria, anxiety, and withdrawal symptoms when access to alcohol is limited. These negative emotional responses have been shown to be primarily mediated by the release of corticotropin releasing factor in the extended amygdala (Koob and Volkow 2010). The uncinate fasciculus (UF) is a hook-shaped white matter tract joining the amygdala and orbitofrontal cortex. This tract is thought to play a key role in emotional responses to events and has been implicated in disorders involving emotional regulation (e.g. anxiety disorders, depression, and conduct disorder) (Taylor et al. 2007; Olson et al. 2015; Damme et al. 2017). UF diffusivity has been shown to be related to levels of anxiety and depression in individuals with heroin addiction (Wong et al. 2015). Microstructural changes in UF have been reported in adolescent cigarette smokers (Zhou et al. 2022) and adolescent binge alcohol drinkers (Jacobus et al. 2013; Luciana et al. 2013).

EF domain and the anterior thalamic radiation

In AUD, the EF domain involves impaired impulse control and persistent and habitual alcohol-seeking behavior despite the desire to avoid negative consequences of alcohol use. The top-down executive control functions affected in AUD are primarily mediated by the prefrontal cortex (Goldstein and Volkow 2011), working in conjunction with thalamic, cerebellar, and parietal cortices (Dosenbach et al. 2008). The anterior thalamic radiation is a major thalamocortical projection between the dorsolateral prefrontal cortex and the dorsomedial thalamic nucleus through the internal capsule's anterior limb and is known to be involved in EF (Mamiya et al. 2018; Niida et al. 2018; Haddad et al. 2021). Studies have linked alterations in ATR microstructure with heightened impulsivity (Joutsa et al. 2011; Yuan et al. 2017; Alfano et al. 2021). Recent evidence suggests that substance use disorders may be associated with decreased ATR volume (Pando-Naude et al. 2021).

White matter microstructure metrics

We assessed white matter microstructure in each of the three pathways using diffusion kurtosis imaging (DKI), a framework

for modeling the diffusion signal that improves upon the standard tensor model by accounting for deviations from Gaussianity (Jensen and Helpern 2010). We focused on anisotropy of both the diffusion tensor (fractional anisotropy, FA) and kurtosis tensor (kurtosis fractional anisotropy, KFA). These metrics range from 0 to 1, where 0 represents perfect isotropy and 1 represents complete directional cohesion along the primary axis.

FA was chosen as the primary metric of interest due to its widespread clinical and research use, allowing for easier comparisons to prior research. KFA was also examined because it has been shown to offer supplementary information and improved contrast in complex tissues and deep brain structures where FA often approaches zero (Hansen and Jespersen 2016). We and others have hypothesized that KFA's enhanced ability to distinguish various microstructural compositions of white matter results in an improvement over FA in diagnostic sensitivity (Hansen and Jespersen 2016; Hansen 2019; Zhang et al. 2019). By including both metrics, we aimed to leverage the more advanced kurtosis model while maintaining the ability to compare our results to prior studies which used the standard tensor model.

To determine whether white matter microstructure is associated with subsequent relapse outcome, we investigated here whether FA and KFA metrics within these three tracts (a) predict dichotomous relapse (relapsed vs. abstained) during a 4-month follow-up period and (b) predict the number of days until relapse during a 4-month follow-up period.

Methods Participants

Participants with AUD were recruited 1–2 weeks following admission to a 28-day in-patient addiction treatment program in Minneapolis, MN as part of a longitudinal study. This analysis focuses on imaging data collected during early abstinence (mean number of days abstinent until MRI session = 25.4, SD = 13.2) and relapse data collected at a four month follow up interview. Written informed consent was obtained from each participant. Participants received monetary compensation for their time. The Institutional Review Board at the University of Minnesota reviewed and approved the consent process and all study procedures. The data that support the findings of this study are available from the corresponding author upon reasonable request.

A total of 86 participants were consented. From these, 11 did not complete the MRI session for the following reasons: 3 left the treatment program and were no longer reachable before the MRI session, 3 participants opted out of the study after consent and before the MRI session, and 5 were found to be no longer eligible (3 because of medical conditions, 1 because of unknown metal in body, and 1 because it was discovered that the identified *primary* substance use disorder diagnosis was not alcohol, but methamphetamine). Of the 75 participants who completed an MRI session, 6 were missing diffusion MRI data due to time constraints during the MRI session. Of the 69 participants from whom we collected diffusion MRI data, 16 were lost to follow-up after the MRI session. As a result, complete data for the scope of this paper (neuroimaging and treatment outcome data) were available for 53 individuals with AUD (Table 1).

Participants underwent random alcohol and drug tests in the treatment program. Each participant's substance use history for the past 6 months before entering the treatment program was recorded using the Timeline Follow-Back (TLFB) (Sobell and Sobell 1992) assessment that was administered for alcohol and for each other substance used (excluding caffeine) (Table 1).

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ne at 4-month follow-up	ated by treatment outcom	Sample separat			
ABS vs. REL	REL $(n = 26)$	ABS $(n = 27)$	All AUD $(n = 53)$		

Table 1. Demographics, history of alcohol use, and clinical self-report measures in individuals with AUD during early abstinence.

Characteristic	All AUD (n=53) Mean or n (SD or %)	ABS $(n = 27)$ Mean or n (SD or %)	REL (n = 26) Mean or n (SD or %)	ABS vs. REL P-value for t-test or χ^2 (italics)
Age	42.8 (10.1)	44.1 (10.8)	41.3 (9.3)	P=0.32
Years of education	14.2 (2.1)	14.3 (2.0)	14.1 (2.2)	P = 0.66
Female, n %	19 (35.9%)	6 (22.2%)	13 (50.0%)	P = 0.09
Employed, n %	21 (39.6%)	12 (44.4%)	9 (34.6%)	P = 0.57
Age of AUD onset	24.7 (10.7)	25.2 (10.8)	24.1 (10.6)	P = 0.70
# of standard drinks: past 6 months	2,799.2 (2,226.8)	2,538.8 (1,876.9)	3,069.5 (2,549.7)	P = 0.39
# drinking days: past 6 months	101.7 (58.1)	101.8 (58.6)	99.5 (58.7)	P = 0.89
# days abstinent before MRI session	22.4 (13.2)	23.5 (10.7)	21.3 (15.5)	P = 0.55
Beck Depression Inventory (BDI) ^a	21.6 (12.0)	21.6 (13.2)	21.7 (10.9)	P = 0.97
State Anxiety Inventory (STAI) ^a	42.2 (11.2)	39.7 (12.0)	44.5 (10.1)	P = 0.12
Penn Alcohol Craving Scale (PACS) ^a	21.9 (7.4)	21.5 (8.4)	22.3 (6.4)	P = 0.71

AUD, alcohol use disorder; MRI, magnetic resonance imaging; SD, standard deviation; ABS, those that remained abstinent in the 4-month follow-up period; REL, those that relapsed during the 4-month follow-up period; χ^2 , chi-square; *p*, significance probability value. ^aOne participant (ABS) did not complete any self-report measures. Four participants (two REL, two ABS) completed PACS and BDI, but not STAI.

Study participants completed clinical self-report measures that have been associated with relapse (Oliva et al. 2018; Stohs et al. 2019; Pareaud et al. 2021) (Table 1): the Beck Depression Inventory (BDI; Beck et al. 1988); the State Anxiety Inventory (STAI; Spielberger et al. 1970); and the Penn Alcohol Craving Scale (PACS; Flannery et al. 1999). Participants completed interviews during a 4-month follow-up period after the neuroimaging session to query dichotomous and continuous relapse metrics.

Dichotomous relapse metrics: group definition

Participants were considered to be in the relapsing group (REL) if they reported consuming at least one drink and/or non-prescribed drug during the 4-month follow-up period. Participants who reported consuming no alcohol or non-prescribed drug during the 4-month follow-up period were considered to be in the abstaining group (ABS).

Continuous relapse metrics

Detailed TLFB (Sobell and Sobell 1992) data were collected from the REL group, recording date of relapse, number of drinks, and number of drinking days after relapse during the 4-month followup period. Time to relapse to alcohol use was operationalized as the number of days until a participant's self-reported first use of alcohol. The variable representing the number of drinks (standard drink, equivalent to 0.6 ounces of pure alcohol (Alcohol Questions and Answers | CDC 2022) after relapse was operationalized as the number of drinks reported to have been consumed in the 4-month follow-up period. The number of days drinking was operationalized as the number of days reported to have consumed at least one drink during the 4-month follow-up period (Table 1).

Imaging data acquisition and analysis

MRI data were collected using a 3 T Siemens Prisma scanner (Siemens, Erlangen, Germany) at the University of Minnesota Center for Magnetic Resonance Research. Acquired images included: a T1-weighted MPRAGE image [TR=2,400 ms, TE=2.24 ms, slices=208, voxel size=0.8 mm³], a T2-weighted SPACE image [TR=3,200 ms, TE=564 ms, slices=208, voxel size=0.8 mm³], and a pair of multi-shell diffusion-weighted images collected using reversed phase-encode blips [TR=3,100 ms, TE=85 ms, slices=92, voxel size=1.5 mm³, multi-band factor=4, *b* values of 0, 1,000, and 2,000 s/mm²].

Image data were preprocessed using the HCP minimal preprocessing pipelines (v4.0.1) (Glasser et al. 2013). For structural data, these steps included alignment of the T1-weighted volume to the T2-weighted volume, correction for gradient distortion and intensity bias, volume segmentation and surface reconstruction using FreeSurfer (v6.0.0) (Dale et al. 1999) and calculation of nonlinear transforms to standard MNI152 space. Diffusion preprocessing included rigid AC-PC alignment to the native structural images, correcting for susceptibility-related distortions using the "topup" tool from the Functional Magnetic Resonance Imaging of the Brain Software Library (FSL; Andersson et al. 2003; Smith et al. 2004), and eddy current distortion with slice outlier replacement using FSL's "eddy" tool (Andersson et al. 2016; Andersson and Sotiropoulos 2016; Andersson et al. 2017; Andersson et al. 2018).

Tractography was carried out using FSL's XTRACT (Warrington et al. 2020), in which expertly defined masks in MNI152 space are warped to native space and used to guide probabilistic tractography of a set of 42 major white matter pathways (19 bilateral and 4 commissural) in each participant, including the 6 tracts of interest (3 bilateral) examined in the present study (Table 2). Tracts of interest were then visually inspected by an experienced rater (D.J.R.) to ensure they were free from anomalies or discernible processing errors. Finally, we refined each tract of interest by excluding voxels with less than 20% of the robust maximum probability of tract membership.

White matter microstructure in each tract was assessed using FA and KFA metrics. FA is a frequently used measure of the anisotropy of the diffusion signal when the tensor model is applied to the data, while KFA reflects the anisotropy of the non-gaussian component (kurtosis tensor) when the higher order kurtosis model is applied to the diffusion signal (Glenn et al. 2015). KFA provides superior contrast in complex tissues compared to standard diffusion tensor models (Hansen and Jespersen 2016). Whole brain FA and KFA scalar maps were calculated from the preprocessed diffusion images using the Diffusion Kurtosis Estimator package (Tabesh et al. 2011). Average tract-wise values were then calculated for each tract, weighted by voxel-wise likelihood of tract membership.

Statistical analysis

First, to determine whether there were significant demographic differences on dichotomous relapse metrics (ABS vs. REL defined

Table 2. Koob–Volkow addiction domains and underlying brain regions and associated white matter pathways.

Domain	Description	Brain Regions (Koob and Volkow 2016; Kwako et al. 2019)	Proposed associated white matter pathway
Reward and Incentive salience (IS)	Exaggerated dopaminergic reward response	Nucleus accumbens Caudate Putamen Globus pallidum Thalamus	Fornix (Fx)
Negative emotionality (NE)	Withdrawal and exaggerated stress when not using	Amygdala Caudate Habenula Nucleus accumbens	Uncinate fasciculus (UF)
Executive functioning (EF)	Preoccupation with drug; heightened impulsivity	Prefrontal cortex Orbitofrontal cortex Hippocampus insula	Anterior thalamic radiation (ATR)

For each white matter pathway, left and right hemispheres were examined separately.

at the 4-month follow-up period), independent samples t-test were conducted on age, years of education, clinical self-report measures (i.e. BDI, STAI, PACS, SCQ), and substance use history (age of onset of alcohol dependence, number of drinks in the past 6 months, number of days drinking in the past 6 months, number of days abstinent until MRI data collection day). To determine whether there were sex (as a biological variable) differences between groups (REL vs ABS), Pearson's chi-square tests were conducted (Table 1).

Binary logistic regression models were fit to assess whether FA and KFA in our tracts-of-interest predicted binary outcome (REL vs ABS) at a 4-month follow-up. Cox proportional hazards models were used to assess whether FA and KFA in our tracts-ofinterest predicted days-to-relapse. For these models, anisotropy metrics were linearly scaled by a factor of 100 to facilitate the interpretability of the coefficient estimates. Each tract-of-interest in each hemisphere was modeled separately and separate analyses were conducted for KFA and FA. Based on the total number of tracts examined (6), a Bonferroni-adjusted significance threshold of 0.0083 was used.

Results

No demographic differences were found between REL and ABS groups (Table 1), Despite there being no significant relationship between sex and age and either continuous or binary relapse measures in our sample, we adjusted for these variables in each model to account for observed trend-level relationships and prior reports of potential effects (Sliedrecht et al. 2019). No significant differences were found in any self-report measures between REL and ABS groups (Table 1).

Diffusion metrics and risk of relapse Dichotomous relapse metric

Descriptive statistics for tract-wise FA and KFA measurements are reported in Table 3. After correcting for the number of comparisons within each diffusion metric by using a Bonferroniadjusted alpha ($\alpha = 0.0083$), only KFA in the right fornix was a significant predictor (P = 0.006) of dichotomous relapse in sex and age-adjusted binary logistic regression models (Table 4); FA in this tract did not survive correction for multiple comparisons (P = 0.018). Results in the bilateral UF suggest that a negative relationship between anisotropy measures and dichotomous relapse may be uncovered in a larger sample, but these did not reach significance in our sample.

Continuous relapse metric

Results of the sex and age-adjusted Cox regression models predicting the number of days to relapse (see Table 5) bore similarities to the logistic regression models predicting dichotomous relapse reported above. In the right fornix, KFA significantly predicted the number of days until relapse (P = 0.002), with increased KFA associated with longer periods of abstinence. FA in this tract showed a similar relationship but did not survive Bonferroni correction (P = 0.011). In the bilateral UF, the relationship between anisotropy measures and abstinence duration was in the same direction as observed in the right fornix but failed to reach significance.

Discussion

This study follows a previous report in which we leveraged the same theory-driven addiction model (Koob and Volkow 2016; Kwako et al. 2019) to examine whether resting state functional connectivity (RSFC) within each addiction domain was associated with binary and continuous relapse metrics (Camchong et al. 2022). This study expands on our brain functional connectivity work by examining whether white matter microstructural alterations in tracts within those same theoretical addiction domains are also predictive of subsequent relapse outcomes (as dichotomous and continuous variables). By examining anatomical metrics that are relatively stable across time, we hoped to both address reliability concerns inherent in RSFC analyses (Noble et al. 2019) and explore potential structural explanations for previously observed RSFC hypoconnectivity.

We examined three pairs of contralateral tracts representing the three addiction domains (Table 2) using two diffusion-based metrics (FA and KFA) that are sensitive to microstructural alterations in white matter. Our approach revealed a relationship between white matter microstructure in one tract associated with the IS domain (right fornix) and non-significant effects in the same direction in the NE domain (bilateral UF). There was no apparent relationship between white matter microstructure and the bilateral ATR, which was selected as representative of the EF domain.

KFA in the right fornix (IS domain) was significantly associated with both dichotomous (relapsed: yes vs. no) and continuous (number of days to relapse) relapse metrics during the 4-month follow-up period, after controlling for age and sex and after correction for multiple comparisons. The IS domain was the

Table 3.	Descriptive	statistics	of tract-wise	FA and KF.	A measurements
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			All AUD		ABS		REL	
Domain	Tract	Hemisphere	FA m (SD)	KFA m (SD)	FA m (SD)	KFA m (SD)	FA m (SD)	KFA m (SD)
EF	Anterior thalamic radiation	Left	0.428 (0.020)	0.502 (0.024)	0.428 (0.020)	0.503 (0.025)	0.427 (0.019)	0.500 (0.025)
IS	Fornix	Left	0.421 (0.018) 0.312 (0.039)	0.307 (0.026)	0.422 (0.020) 0.316 (0.044)	0.312 (0.023)	0.419 (0.017) 0.309 (0.034)	0.302 (0.027)
NE	Uncinate fasciculus	Right Left Right	0.301 (0.042) 0.415 (0.024) 0.416 (0.024)	0.314 (0.049) 0.506 (0.028) 0.507 (0.028)	0.320 (0.035) 0.421 (0.024) 0.423 (0.022)	0.328 (0.043) 0.510 (0.029) 0.516 (0.027)	0.299 (0.046) 0.409 (0.023) 0.408 (0.024)	0.300 (0.051) 0.501 (0.026) 0.498 (0.028)

Table 4.	Predicting	dichotomous r	elanse l	based o	on diffusior	metrics	measured	during	Parl	, abstinence
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			FA			KFA	KFA		
Domain	Tract	Hemisphere	Est.	CI	р	Est.	CI	р	
EF	Anterior thalamic radiation	Left	1.00	(0.74–1.36)	0.996	0.96	(0.74–1.23)	0.763	
		Right	1.01	(0.73-1.40)	0.975	1.04	(0.81-1.33)	0.731	
IS	Fornix	Left	0.94	(0.79-1.11)	0.462	0.93	(0.80-1.07)	0.314	
		Right	0.80	(0.66–0.95)	0.018 ^a	0.79	(0.66–0.92)	0.006 ^b	
NE	Uncinate Fasciculus	Left	0.83	(0.63-1.08)	0.178	0.85	(0.66-1.08)	0.195	
		Right	0.82	(0.62–1.06)	0.140	0.82	(0.64–1.02)	0.085	

^aSignificant only before Bonferroni correction ^bSignificant after correction for multiple comparisons

Table 5.	Predicting	days-to-rela	pse based (on diffusion	metrics me	asured during	g early	v abstinence.

			FA			KFA	KFA		
Domain	Tract	Hemisphere	Est.	CI	р	Est.	CI	р	
EF	Anterior thalamic radiation	Left	0.95	(0.77-1.17)	0.628	0.94	(0.79–1.12)	0.476	
		Right	0.98	(0.78-1.23)	0.861	1.02	(0.86-1.21)	0.818	
IS	Fornix	Left	0.98	(0.87-1.10)	0.707	0.96	(0.88-1.06)	0.453	
		Right	0.86	(0.76–0.96)	0.011 ^a	0.86	(0.78–0.95)	0.002^{b}	
NE	Uncinate fasciculus	Left	0.86	(0.71-1.05)	0.137	0.90	(0.77-1.05)	0.170	
111		Right	0.85	(0.71–1.01)	0.064	0.86	(0.74–1.01)	0.062	

^aSignificant only before Bonferroni correction ^bSignificant after correction for multiple comparisons

only domain in which a significant relationship was found with either dichotomous or continuous relapse. These findings align with our prior results based on RSFC strength, where IS domain hypoconnectivity was predictive of both continuous and dichotomous relapse outcomes and was the only domain that remained a significant predictor of dichotomous relapse after controlling for sex (Camchong et al. 2022). Due to its role in the mesolimbic reward pathway, microstructural abnormalities in the fornix may be associated with cue-induced alcohol craving and habit-like compulsive alcohol consumption (Koob and Volkow 2016).

Though not significant, increased FA and KFA in the bilateral UF (NE domain) were associated with decreased odds of continuous and dichotomous relapse in our sample. Microstructural abnormalities in the UF likely influence emotional processing (Von Der Heide et al. 2013) which could exacerbate symptoms of protracted withdrawal in those undergoing treatment for AUD (Koob and Volkow 2016). Although not significant in our limited sample, the directionality and strength of the observed relationship suggest that further exploration of this relationship is merited.

In the IS and NE domains, the direction of all observed relationships was negative, with increases in anisotropy corresponding to decreased risk of relapse and longer periods of sustained abstinence and vice versa. Under the common interpretation of anisotropy as a proxy for white matter integrity, this is the expected directionality and suggests that relapse risk in AUD may be related to white matter that is damaged or underdeveloped. Impaired structural connectivity may also explain our previous observation of decreased RSFC in these domains (Camchong et al. 2022). However, caution must be taken in interpreting diffusion metrics and further exploration of these findings is warranted (see *Considerations*).

Comparison to prior research

Several previous studies have also investigated white matter microstructure in the context of relapse in AUD. Using a design similar to the present study, Zou et al. (2018) reported decreased FA in the genu and splenium of the corpus callosum as well as the right fornix/stria terminalis in participants who went on to relapse compared to those who remained abstinent. Sorg et al. (2012) also reported lower FA in callosum, bilateral UF, right superior corona radiata, and left anterior internal capsule in participants who returned to heavy use compared to those who abstained as measured six months after treatment. These studies used a whole brain, tract-based spatial statistics (TBSS) approach that substantially differs from our tractography-driven approach. Nonetheless, our observations While the present study focused on white matter microstructure as a predictor of relapse in AUD, it is important to also consider how microstructure is itself impacted by abstinence/relapse. In a relevant longitudinal study, Pfefferbaum et al. (2014) reported that age-related decreases in FA across wide areas of the brain were exaggerated in participants with AUD who relapsed after treatment, while those who abstained showed a flatter trajectory suggesting that FA may recover to normal ranges in prolonged abstinence. Several other studies have also presented evidence that white matter alterations in AUD may be at least partially reversible (Gazdzinski et al. 2010; Alhassoon et al. 2012; Zou et al. 2017).

Considerations

We relied on a theoretical model of addiction (Koob and Volkow 2010; Koob and Volkow 2016) to inform the a priori selection of several white matter tracts in which we hypothesized that microstructural measures may relate to relapse risk in AUD. In the context of our limited sample size, we lack sufficient power to address the *specificity* of our findings and thus cannot determine whether similar effects may also be found elsewhere in the brain. Furthermore, the theoretical model on which we relied was derived in large part from research on animal models of addiction that may not necessarily extend to AUD in humans. Although this analysis leveraged the Koob–Volkow model in its design, the results should not be interpreted as supporting or refuting the model's validity.

Although diffusion MRI is a useful tool for examining white matter microstructure in vivo, caution must be taken when interpreting how these metrics may reflect the biophysical properties of the underlying tissue (Mueller et al. 2015). A variety of microstructural features may influence the diffusion signal and therefore metrics derived from it, including myelination, axonal damage, and orientation of crossing fibers (De Erausquin and Alba-Ferrara 2013). Our primary goal was to assess whether measurable microstructural alterations were associated with relapse in AUD. The *nature* of those differences is beyond the scope of this study. Furthermore, we are unable to infer from our data whether variations in FA and KFA were caused by chronic alcohol use or if they were pre-existing, perhaps even predisposing individuals to AUD. These questions are well suited for future research.

Interpretation of these findings in relation to our recent report on RSFC in Koob–Volkow domains (Camchong et al. 2022) requires some additional considerations. The white matter tracts chosen to represent each domain in the present study were selected from the 23 expertly defined pathways available in FSL's XTRACT. Perfect correspondence, wherein each pathway connects those structures and only those structures that comprised the RSFC networks used in our prior study, was not feasible. Further investigations using alternative tractography approaches may provide additional information. Furthermore, we do not provide direct evidence to determine whether the observed microstructural effects cause resting state hypoconnectivity or instead reflect structural degradation as a result of sustained hypoconnectivity.

Conclusions

We observed that decreases in anisotropy measures of the right fornix, a white matter tract involved in the IS domain, were related to an increased risk of dichotomous relapse as well as shorter duration of abstinence among individuals with AUD in an inpatient treatment program. No significant relationships between microstructure and relapse outcomes were found in tracts representing the EF and NE domains, but the observed relationship between microstructure and relapse in NE domain tracts (bilateral UF) encourages further exploration using a larger sample. This study supplements our previous report in which reduced RSFC in domain-specific networks was associated with increased relapse risk in AUD. The current findings suggest that some microstructural features may also mediate relapse risk. Further investigation is needed into how the structure and function of these addiction networks may modulate resilience to relapse.

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