

## RESEARCH ARTICLE

# Shifted balance of dorsal versus ventral striatal communication with frontal reward and regulatory regions in cannabis-dependent males

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## Abstract

The transition from voluntary to addictive behavior is characterized by a loss of regulatory control in favor of reward driven behavior. Animal models indicate that this process is neurally underpinned by a shift in ventral–dorsal striatal control of behavior; however, this shift has not been directly examined in humans. The present resting state functional magnetic resonance imaging (fMRI) study employed a two-step approach to: (a) precisely map striatal alterations using a novel, data-driven network classification strategy combining intrinsic connectivity contrast with multivoxel pattern analysis and, (b) to determine whether a ventral to dorsal striatal shift in connectivity with reward and regulatory control regions can be observed in abstinent (28 days) male cannabis-dependent individuals ( $n = 24$ ) relative to matched controls ( $n = 28$ ). Network classification revealed that the groups can be reliably discriminated by global connectivity profiles of two striatal regions that mapped onto the ventral (nucleus accumbens) and dorsal striatum (caudate). Subsequent functional connectivity analysis demonstrated a relative shift between ventral and dorsal striatal communication with fronto-limbic regions that have been consistently involved in reward processing (rostral anterior cingulate cortex [ACC]) and executive/regulatory functions (dorsomedial prefrontal cortex [PFC]). Specifically, in the cannabis-dependent subjects, connectivity between the ventral striatum with the rostral ACC increased, whereas both striatal regions were uncoupled from the regulatory dorsomedial PFC. Together, these findings suggest a shift in the balance between dorsal and ventral striatal control in cannabis dependence. Similar changes have been observed in animal models and may promote the loss of control central to addictive behavior.

## KEYWORDS

anterior cingulate, addiction, cannabis, cognitive control, intrinsic connectivity contrast, data-driven, functional connectivity, prefrontal cortex, reward, striatum

## 1 | INTRODUCTION

Converging evidence from animal and human research indicates that maladaptations in fronto-limbic-striatal circuitries drive the loss of control that characterizes addiction (Fineberg et al., 2010; Morein-Zamir &

Robbins, 2015). The striatum lies at the core of this circuitry and critically contributes to both, acute drug reinforcement and the transition from voluntary to addictive use that is accompanied by a loss of control (Brand, Young, Laier, Wölfling, & Potenza, 2016; Everitt & Robbins, 2016; Fineberg et al., 2010).

The striatum contributes to several domains that undergo critical adaptations during the transition to addiction, including associative

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learning, behavioral control, incentive salience, and reward (Haber, 2016; Robbins, Gillan, Smith, de Wit, & Ersche, 2012). The functional heterogeneity of the striatum and its multifaceted contributions to addiction are mirrored in its complex organization into distinct subregions that communicate with the entire cortex via subregion-specific fronto-striatal loops (Haber, 2016). Based on their specific functions and projections, the striatum is divided into the ventral striatum (VS) that exhibits strong connections with ventral anterior cingulate and orbitofrontal regions regulating reward and reinforcement behavior, and the dorsal striatum (DS) that exhibits strong connections with dorsolateral and dorsomedial prefrontal regions engaged in regulatory control (Di Martino et al., 2008; Haber, 2016; Postuma & Dagher, 2006). This functional and network-level segregation has important implications for the transition to addiction, with overarching conceptualizations proposing that the loss of control is accompanied by a progressive shift from controlled toward reward-driven impulsive behavior (Jentsch & Taylor, 1999; Noël, Brevers, & Bechara, 2013).

Partly driven by the ongoing discussion about cannabis legalization and concomitantly increasing demand for treatment of cannabis dependence (EMCDDA, 2008; Volkow et al., 2016), a number of studies have examined effects of cannabis use on the brain (overview see, e.g., Lorenzetti, Solowij, & Yücel, 2016; Weinstein, Livny, & Weizman, 2016). In line with the functions of the striato-frontal circuitry and its proposed contribution to addiction task-based neuroimaging studies demonstrated exaggerated striatal reactivity to cannabis cues (Filbey et al., 2016), reward anticipation (Jager, Block, Luijten, & Ramsey, 2013) and decreased frontal activity during executive control (Weinstein et al., 2016; Wrege et al., 2014; Yanes et al., 2018) in chronic users. This pattern may reflect exaggerated striatal reactivity in the context of deficient prefrontal control and mirror maladaptations that have been observed across addictive disorders (Brand et al., 2016; Goldstein & Volkow, 2011).

Despite the important contributions of these studies to characterize mechanisms of cannabis dependence, the task-based approach does not allow direct evaluation of the ventral to dorsal striatal shift due to: (a) limitation of task-based fMRI to regions engaged by the experimental paradigm (Weinstein et al., 2016) and (b) stimulus and context dependences of striatal alterations in cannabis dependence (Gilman, 2017; Zimmermann et al., 2017). Resting state fMRI functional connectivity (rsfMRI-FC) approaches allow a more holistic assessment of functional alterations in the absence of task or contextual modulation. Moreover, these approaches have a high sensitivity for transient and lasting striatal alterations on the subregion-specific network level (Di Martino et al., 2008; Di Martino et al., 2011). Two recent studies applied rsfMRI-FC to determine lasting disruptions in the frontostriatal functional circuitries in cannabis dependence (Blanco-Hinojo et al., 2017; Zimmermann, Yao, et al., 2017). However, findings were limited by the confirmatory nature of the rsfMRI-FC analysis (Zimmermann, Yao, et al., 2017) or the use of literature-based striatal seed regions (Blanco-Hinojo et al., 2017) that have a high sensitivity to capture subregion specific functional networks in healthy subjects (Di Martino et al., 2008) but might not be sensitive to the specific alterations related to cannabis dependence. Despite supporting evidence for a striatal shift in response to masked drug cues in cannabis dependence (Wetherill et al., 2016), the ventral to dorsal

shift has not been explicitly examined in the absence of external stimulation.

The present study therefore employed a two step approach to: (a) precisely map striatal alterations in cannabis-dependent individuals using network classification and (b) determine whether a ventral to dorsal striatal shift in connectivity with reward and regulatory control regions can be observed. In an initial step, intrinsic striatal alterations were mapped using a novel fully data-driven network classification approach that operates independently of a priori assumptions about striatal architecture and thus enables higher sensitivity to detect striatal functional changes (Martuzzi et al., 2011; Rubinov & Sporns, 2010). Next, altered functional communication of the determined striatal subregions was examined on the whole-brain network level (for a similar approach see Walpola et al., 2017). To this end, rsfMRI data were acquired in  $n = 28$  participants that fulfilled the diagnostic and statistical manual of mental disorders, fourth edition (DSM-IV) criteria for cannabis dependence and  $n = 28$  matched healthy controls. To control for subacute effects of cannabinoid metabolites (Vandevenne, Vandebussche, & Verstraete, 2000), craving (Copersino et al., 2006), and rapid neural recovery of homeostatic receptor adaptations after cessation of chronic cannabis use (Hirvonen et al., 2012) participants underwent 28 days of cannabis abstinence before acquisition of the MRI data.

Based on current neurobiological frameworks suggesting that the transition to addiction is accompanied by regional-specific neuroplastic changes in the ventral and dorsal striatum and accumulating evidence from task-based neuroimaging studies suggesting that chronic cannabis use is associated with aberrant reward and regulatory control processes (Everitt & Robbins, 2016; Weinstein et al., 2016; Wrege et al., 2014; Yanes et al., 2018) we expected to observe corresponding maladaptations on the level of the network level organization of the brain. Specifically, we hypothesized that: (a) global connectivity differences between the groups specifically map to the DS and VS, reflecting regional-specific striatal adaptations and (b) a shift in the network communication of the striatal subregions reflecting increased communication within frontostriatal pathways related to reward processing and concomitantly decreased communication in pathways related to cognitive control.

## 2 | METHODS

### 2.1 | Participants

For the present study, data from  $n = 16$  male cannabis-dependent users and  $n = 16$  healthy male controls from our previous publication on convergent task and resting functional connectivity alterations in cannabis users was included (Zimmermann, Yao, et al., 2017). To enhance statistical power for the present data-driven approach, the sample size was increased to  $n = 28$  male cannabis-dependent subjects and  $n = 28$  nonusing controls using identical inclusion and exclusion criteria. To control for potential confounding influences of menstrual cycle on striatal resting state activity as well as complex interactions between menstrual cycle and addiction-related alterations in frontostriatal functioning (Franklin et al., 2015; Wetherill,

Jagannathan, Hager, Maron, & Franklin, 2016; Wiers et al., 2016), only male subjects were included (for a similar approach see Zimmermann, Walz, et al., 2017).

All cannabis users were diagnosed with a current cannabis dependence according to DSM-IV during the 18 months before the examination. Diagnostic criteria were assessed by the structured clinical interview (SCID for DSM-IV) conducted by a trained psychologist. To facilitate the determination of long-term alterations associated with cannabis dependence that persist beyond subacute cannabinoid metabolites (Vandevenne et al., 2000), craving (Copersino et al., 2006), and rapid neural recovery (Hirvonen et al., 2012) participants were required to abstain from cannabis for 28 days prior to the assessment. Comparable abstinence periods have been employed in previous studies to differentiate enduring (long term) effects of cannabis use from transient (subacute) alterations and rapid neural recovery (overview in Crean, Crane, and Mason (2011); Ganzer, Bröning, Kraft, Sack, and Thomasius (2016); Schreiner and Dunn (2012)).

At the time point of recruitment, most cannabis-dependent subjects were active cannabis users or in early phases of abstinence. Inclusion criteria for all participants were: (a) age between 18 and 35 years, (b) right-handedness, (c) negative urine toxicology for cannabis and other illicit drugs (immunoassay, substance/cut-off per milliliter: THC/50 ng, amphetamines/500 ng, cocaine/300 ng, methamphetamine/500 ng, MDMA/300 ng, opiates/300 ng, methadone/300 ng), and (d) self-reported abstinence of >28 from cannabis validated by negative urine screens on the day of the fMRI assessment. One user reported having used cannabis on one occasion 14 days before the experiment, but was included due to negative urine toxicology. Exclusion criteria for all participants were: (a) profound DSM-IV axis I or axis II disorders, such as psychotic or bipolar symptoms, (b) Beck Depression Inventory score > 20 indicating a moderate depression, (c) medical disorder, (d) current/regular medication, and (e) use of other illicit substances >75 lifetime occasions. Controls were included if their cumulative lifetime use of cannabis was <15 g.

Anxiety, mood and concentration were assessed on the day of the examination using validated scales including the social interaction anxiety scale (SIAS) (Mattick & Clarke, 1998), the state-trait anxiety inventory (STAI) (Spielberger, Gorsuch, Lushene, Vagg, & Jacobs, 1983), positive and negative affect schedule (Crawford & Henry, 2004), and the d2 test of attention (Brickenkamp & Zillmer, 1998). These measures allowed control for potential confounding effects of common withdrawal symptoms. Cannabis use patterns and the use of other drugs were assessed using corresponding modules of a structured interview that has been used in previous studies of our group (e.g., Becker et al., 2015; Becker, Wagner, Gouzoulis-Mayfrank, Spuentrup, & Daumann, 2010; Zimmermann, Yao, et al., 2017). Differences in potential confounders were examined by independent sample *t* test; in case of a nonnormal distribution of the underlying data nonparametric analyses were used. As a trade-off between effects of acute nicotine and nicotine craving on neural activity all participants underwent a supervised abstinence from nicotine 1.5 hr before the fMRI assessment. Cannabis-dependent subjects were recruited in cooperation with the LVR Clinics Bonn, Germany, Addiction Department that offers a specialized treatment program for cannabis

dependence. Potential participants were contacted and invited for an initial screening session to further validate study participation criteria. Participants who agreed to stay abstinent for at least 28 days were scheduled for the fMRI assessment on a separate day following the abstinence period. Controls were recruited by advertisements and underwent identical study procedures. Written informed consent was obtained and study protocols were approved by the local ethics committee (Medical Faculty, University of Bonn).

Three cannabis users were excluded due to a history of exceeding cause of other illicit drugs and one was excluded due to excessive head movement during fMRI, leading to a final sample of  $n = 24$  cannabis users and  $n = 28$  healthy controls.

## 2.2 | MRI data acquisition

Functional and structural MRI was acquired using a Siemens TRIO 3-Tesla system with a 32-channel head coil. rsfMRI data were acquired before any tasks using a T2\*-weighted echo-planar imaging (EPI) pulse sequence (repetition time = 2,580 ms, echo time = 30 ms, number of slices = 47, slice thickness = 3.5 mm, no gap, field-of-view =  $224 \times 224 \text{ mm}^2$ , resolution =  $64 \times 64$ , flip angle =  $80^\circ$ , number of volumes = 180). To improve spatial normalization and exclude participants with apparent brain pathologies, a high-resolution whole-brain volume T1-weighted images was acquired in addition using a 3D spoiled gradient echo pulse sequence (repetition time = 1,660 ms, echo time = 2.54 ms, flip angle =  $9^\circ$ , field-of-view =  $256 \times 256 \text{ mm}^2$ , acquisition matrix =  $256 \times 256$ , thickness = 0.8 mm, number of slices = 208).

Participants were instructed to lie still and relax with their eyes closed while thinking of nothing in particular, yet not to fall asleep during the resting state acquisition.

## 2.3 | Preprocessing of the functional resting state data

rsfMRI data were preprocessed using standard statistical parametric mapping (SPM), analysis of functional neuroImages (AFNI) and FMRIB software library (FSL) routines in combination with advanced independent component analysis (ICA-AROMA, Pruim et al., 2015). The first five volumes were discarded to allow MRI T1 equilibration. The remaining volumes were slice-time corrected, spatially realigned to the first volume, and unwarped to correct for nonlinear distortions possibly related to head motion or magnetic field inhomogeneity using SPM12 (<http://www.fil.ion.ucl.ac.uk/spm/software/spm12/>). The functional time series were further processed using FSL (<http://www.fmrib.ox.ac.uk/fsl>), including nonbrain removal using BET (Smith, 2002), spatial smoothing with a Gaussian kernel of 6 mm full width at half maximum, and grand-mean scaling. Subsequently, the data were submitted to an ICA for automatic removal of motion artifacts (ICA-AROMA, Pruim, Mennes, van Rooij, et al., 2015)—a procedure that has been shown to minimize the impact of motion on functional connectivity metrics and decrease the loss in temporal degrees of freedom compared to spike regression and scrubbing (Pruim, Mennes, Buitelaar, & Beckmann, 2015). Next, mean signals from white matter (WM) and cerebrospinal fluid (CSF), linear and quadratic drift across

time, together with a series of sine and cosine functions removing all frequencies outside the range (0.01–0.08 Hz), were regressed out in a single regression step using AFNI's 3dTproject. Similar to the preprocessing protocol used by Pruim, Mennes, van Rooij, et al. (2015), WM and CSF time series were derived by determining the mean time series over voxels within predefined subject-specific WM and CSF masks. To obtain these masks, we first thresholded the MNI152 average CSF and WM priors maps (95% of the robust range) and subsequently registered to native EPI space. Likewise, we applied FSL FAST (Zhang, Brady, & Smith, 2001) to the individual T1 images to derive a CSF and WM probability map and thresholded at 95% followed by registration to native EPI space. Multiplication of both masks resulted in the respective conservative CSF and WM masks. Finally, functional MRI data were registered to T1 and standard MNI152 space using Boundary-Based Registration as implemented in FLIRT (Jenkinson, Bannister, Brady, & Smith, 2002; Jenkinson & Smith, 2001) and FNIRT nonlinear registration (Andersson, Jenkinson, & Smith, 2010) with 6 and 12 df, respectively, and interpolated to  $2 \times 2 \times 2$  mm.

## 2.4 | Quality control for motion artifacts

One participant (cannabis user) was excluded due to excessive head movement ( $>2.5$  mm or  $2.5^\circ$  absolute motion over the whole scan). Mean-frame-wise displacement (FD, Power, Barnes, Snyder, Schlaggar, & Petersen, 2012) and spike control was employed for data quality assurance and motion control. FD was calculated with the raw functional data for the remaining participants. Importantly, the two groups (controls and cannabis users) did not differ in mean FD (controls: mean  $\pm$  SD =  $0.087 \pm 0.032$ , range from 0.044 to 0.145; cannabis users: mean  $\pm$  SD =  $0.083 \pm 0.024$ , range from 0.042 to 0.142,  $p = .568$ , Cohen's  $d = 0.160$ ). We also defined a spike in head movements as FD  $> 0.3$  mm. Less than 7% scans were regarded as spikes in all of the participants and the two groups did not differ in the number of spikes (control: mean  $\pm$  SD =  $2.427 \pm 3.501$ , range from 0 to 11; user: mean  $\pm$  SD =  $1.750 \pm 2.723$ , range from 0 to 11.  $p = .445$ , Cohen's  $d = 0.214$ ).

## 2.5 | Determining altered striatal connectivity in cannabis dependence using the intrinsic connectivity contrast

To promote an unbiased determination of striatal subregions exhibiting altered functional connectivity related to cannabis dependence, a data-driven network level approach was implemented (Walpola et al., 2017). To this end, whole-brain voxel-to-voxel connectivity was computed for each striatal voxel using the intrinsic connectivity contrast (ICC), an index similar to "degree" in graph theory but without the need for a correlation threshold, which reflected the average  $r^2$  of a given voxel with all other brain voxels (Martuzzi et al., 2011; Rubinov & Sporns, 2010). The corresponding ICC analysis was implemented by weighting the connections of a given voxel with every other voxel in the brain (restricted to the SPM gray matter mask  $>0.3$ ) by their  $r^2$  value (Martuzzi et al., 2011), thus both negative and positive correlations will be captured in positive ICC values.

Next, resulting ICC maps were used for discriminating between cannabis-dependent subjects and healthy controls using a multivoxel

pattern analysis (MVPA) with a linear kernel support vector classifier as implemented in LIBSVM ([www.csie.ntu.edu.tw/~cjlin/libsvm](http://www.csie.ntu.edu.tw/~cjlin/libsvm)). A searchlight procedure with a three-voxel radius was used to provide measures of classification accuracy in the neighborhood of each voxel in the striatum mask which was obtained from the Harvard-Oxford Subcortical Structural Atlases (subcortical HO atlas, Harvard Center for Morphometric Analysis) by combining the accumbens, caudate, putamen, and pallidum masks (thresholded at 0% population probability) that were further masked by the gray matter mask to remove voxels with low gray matter signal. Classification was evaluated by a 10-fold cross validation during which all participants were randomly assigned to 10 subsamples of five or six participants using MATLAB's `cvpartition` function. In each iteration, the optimal hyperplane was computed based on the multivariate pattern of ICC values of 47 or 46 participants and evaluated by the excluded five or six participants. The default cost parameter ( $C = 1$ ) was automatically corrected for unbalanced groups by adjusting weights inversely proportional to group frequencies in the training data. The iteration was repeated 10 times with each group being the testing set once and then the average classification accuracy of each sphere was assigned to the center voxel in the sphere. The training set was scaled to  $[-1, 1]$ , and the testing set was scaled using the same scaling factors before applying SVM (Hsu, Chang, & Lin, 2003). To avoid a potential bias of train-test splits, the cross-validation procedure was repeated five times by producing different splits in each repetition and the resultant maps were averaged to produce a convergent estimation. To test whether the resultant measures exceeded chance level, we used permutation tests to simulate the probability distribution of the classification. Briefly, we randomly shuffled the group labels and recomputed these measures (repeated 5,000 times) to build empirical distributions. The resultant maps were then converted to  $p$  values and familywise error (FWE) corrected for multiple comparisons.

## 2.6 | Follow-up seed-to-voxel functional connectivity

Given that ICC is predominantly an exploratory strategy that allows a data-driven precise mapping of striatal regions characterized by their functional connectivity profile differences between the groups functional connectivity analysis was used to determine the associated networks that contribute to these differences (for a similar approach see Martuzzi et al., 2011; Walpola et al., 2017). Importantly, the rationale for the present analyses was based on addiction animal models that indicate a ventral-to-dorsal shift in striatal processing underlying the development of addiction (Everitt & Robbins, 2016). In line with our expectations, the MVPA on the ICC revealed group differences in the global connectivity profiles of ventral and dorsal striatal regions (details see Section 3). Consequently, the major aim of the follow-up functional connectivity analysis was to determine regions that shift communication with ventral vs dorsal striatal nodes in drug dependence. To this end, two 6-mm-radius spheres (masked by the striatum mask and the gray matter mask) centered at the peak voxels derived from the MVPA results (DS and VS) were used as seed regions, and Fisher  $z$ -transformed image maps were then created by calculating the correlation coefficient of each voxel in the brain with the mean

**TABLE 1** Group characteristics and drug use parameters

Measure	Cannabis users mean (SD)	Controls mean (SD)	p
Age (years)	24.00 (3.46)	23.39 (2.86)	.49
Education (years)	14.50 (11.00–22.00) <sup>a</sup>	14.25 (12.00–19.00) <sup>a</sup>	.75 <sup>b</sup>
d2 concentration performance	196.46 (39.61)	205.43 (50.57)	.49
STAI-state	34.17 (7.70)	31.07 (7.71)	.16
STAI-trait	35.87 (8.40)	32.64 (7.22)	.14
SIAS	16.50 (8.00–46.00) <sup>a</sup>	16.00 (5.00–33.00) <sup>a</sup>	.24 <sup>b</sup>
Number of smokers	N = 22	N = 23	
Years of nicotine use	8.23 (4.49)	6.24 (3.72)	.11
Cigarettes per day	8.00 (2–20.00) <sup>a</sup>	8.00 (1–30.00) <sup>a</sup>	.96 <sup>b</sup>
Pack-year	4.16 (3.59)	3.42 (3.84)	.50
Alcohol occasions per week	1.00 (0–4.00) <sup>a</sup>	1.00 (0–4.00) <sup>a</sup>	.74 <sup>b</sup>
Alcohol units per week	5.00 (0–24.00) <sup>a</sup>	4.90 (0–18.00) <sup>a</sup>	.92 <sup>b</sup>
PANAS-positive	33.00 (16.00–42.00) <sup>a</sup>	32.50 (22.00–48.00)	.67 <sup>b</sup>
PANAS-negative	11.00 (10.00–25.00) <sup>a</sup>	10.00 (10.00–14.00) <sup>a</sup>	.10 <sup>b</sup>
Past ecstasy use	N = 15	N = 2	–
Lifetime occasions ecstasy	10.00 (1–75) <sup>a</sup>	(1, 8)	
Past cocaine use	N = 11	N = 0	–
Lifetime occasions cocaine	6.00 (1–70) <sup>a</sup>	–	
Past amphetamine use	N = 15	N = 1	–
Lifetime occasions amphetamine	20.00 (1–75) <sup>a</sup>	30.00	
Past hallucinogen use	N = 9	N = 0	–
Lifetime amount hallucinogen	6.00 (1–50) <sup>a</sup>	–	
Past opiate use	N = 3	N = 1	–
Lifetime occasions opiate	(1, 1, 4)	30.00 <sup>c</sup>	
Past cannabis use	N = 24	N = 22	–
% lifetime cannabis dependence	100%	0%	

Abbreviations: PANAS = positive and negative affect schedule; SIAS = social interaction anxiety scale; STAI = state–trait anxiety inventory.

<sup>a</sup> Median (range).

<sup>b</sup> Mann–Whitney *U* test.

<sup>c</sup> Prescription medicinal use.

time-series of the seed regions. To determine the shift in ventral vs dorsal striatal networks associated with dependence, the subject-level seed-specific *Z*-maps were submitted to a two-way mixed ANOVA with seed (VS and DS) entered as a within-subject factor and group (controls and cannabis users) entered as a between-subject factor employed in the FSL's Randomise Tool. Corrections for multiple comparisons were conducted using permutation-based inferences (10,000 permutations) with Threshold-Free Clustering Enhancement (TFCE) which provides a strict control while improving replicability (Chen, Lu, & Yan, 2018; Smith & Nichols, 2009). For regions showing significant interaction effects (*F* test), the mean Fisher *z*-transformed correlation coefficients were extracted from the underlying anatomical regions for further post hoc analysis within SPSS.

## 2.7 | Validation using literature-based VS and DS seeds

In addition, to validate the seed-to-voxel functional connectivity results, functional connectivity analyses were then rerun using predefined VS and DS masks. An anatomical VS mask was defined as nucleus accumbens (subcortical HO atlas) with the probability

threshold set to 30% (228 voxels in total). Following Tinnermann, Geuter, Sprenger, Finsterbusch, and Büchel (2017), coordinates from different studies (Daw, O'doherty, Dayan, Seymour, & Dolan, 2006; Fitzgerald et al., 2011; Luijten, Schellekens, Kühn, Machielse, & Sescousse, 2017; Schönberg, Daw, Joel, & O'Doherty, 2007; Sweitzer et al., 2016) were averaged to create the mean coordinates of reported DS peak voxels. For those lateralized coordinates we multiplied *x* coordinates by  $-1$  to cover both hemispheres. The DS mask was then created by drawing two 6-mm-radius spheres centered at  $\pm 13, 11, 14$  (for bilateral DS) and masked by the striatum mask (223 voxels in total). To control for nongray matter signal, including WM and CSF, both VS and DS masks were masked by the gray matter mask.

## 2.8 | Functional characterization of the determined networks

To support the functional characterization of the networks that exhibit a dependence-associated shift in VS versus DS communication, the NeuroSynth decoder function (<http://neurosynth.org/decode/>) was used to employ a large-scale automated meta-analysis map. The top 25 terms (excluding terms for brain regions) ranked by

**TABLE 2** Cannabis use parameters

Cannabis use parameter	Cannabis users mean $\pm$ SD (range)	Controls mean $\pm$ SD (range)
N	24	22
Age of first cannabis use	15.17 $\pm$ 1.23 (13–17)	16.36 $\pm$ 1.58 (13–19)
Days since last cannabis use	30.00 <sup>a</sup> (14–500)	670.00 <sup>a</sup> (120–3,000)
Maximum frequency of cannabis use (days/month)	28.08 $\pm$ 4.51 (14–30)	–
Duration of regular cannabis use (months)	78.38 $\pm$ 36.40 (19–144)	–
Lifetime amount of cannabis (g)	1,616.5 <sup>a</sup> (62–5,786)	2 <sup>a</sup> (0.1–12)

<sup>a</sup> Median.

the correlation strengths between the regions exhibiting dependence-associated alterations and the meta-analytic map were visualized using word cloud with the size of the font scaled by correlation strength.

### 3 | RESULTS

#### 3.1 | Group characteristics

The groups were comparable with respect to important potential confounders including socio-demographics, attention and the use of licit drugs (Table 1). The lack of significant group differences in attention, mood, and state anxiety (Table 1) argues against strong confounding effects of cannabis withdrawal. As expected cannabis users reported greater lifetime experiences with other illicit drugs (Table 1) than controls. Cannabis use parameters are reported in Table 2. In post-MRI interviews, none of the subjects reported having fallen asleep during the scan.

#### 3.2 | Intrinsic connectivity contrast

The searchlight MVPA identified brain regions within the striatum that exhibit altered global connectivity patterns in cannabis users compared to controls. Specifically, the groups could be reliably discriminated in two regions that mapped onto the VS (peak voxel coordinates, 10, 12, –10 mapping onto the right nucleus accumbens, accuracy = 84.62%, voxel-wise FWE-corrected  $p = .040$ ) and the DS (peak voxel coordinates, –12, 14, 16 mapping onto the left caudate nucleus, accuracy = 85.38%, voxel-wise FWE-corrected  $p = .026$ ) (Figure 1).

#### 3.3 | Follow-up analyses—seed-to-voxel functional connectivity

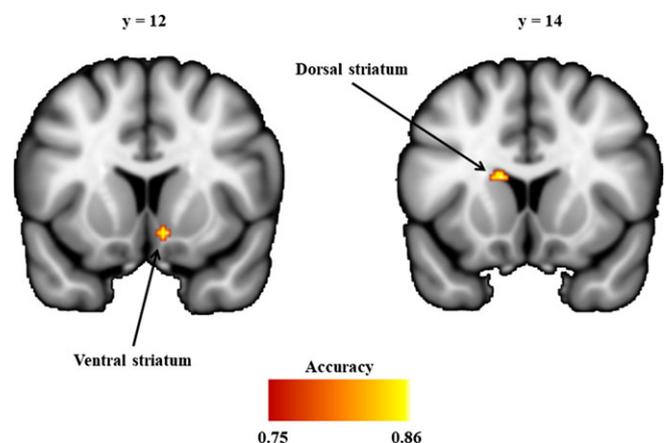
The main effects of seed reflecting the differential connectivity patterns of the VS versus DS seeds further confirmed the mapping of the MVPA to the VS and DS: (a) the VS seed showed stronger connectivity with ventral regions of the frontal cortex including rostral anterior

cingulate cortex (rACC) and orbitofrontal regions, whereas the DS seed exhibited stronger connectivity with the dorsal frontal regions including dorsolateral prefrontal cortex (dmPFC), a pattern resembling previous human and animal research (Di Martino et al., 2008; Haber, 2016). (b) There was a strong overlap between the networks identified for the VS and DS regions determined in the present study and the literature-based seeds from the validation analysis (Figure 2).

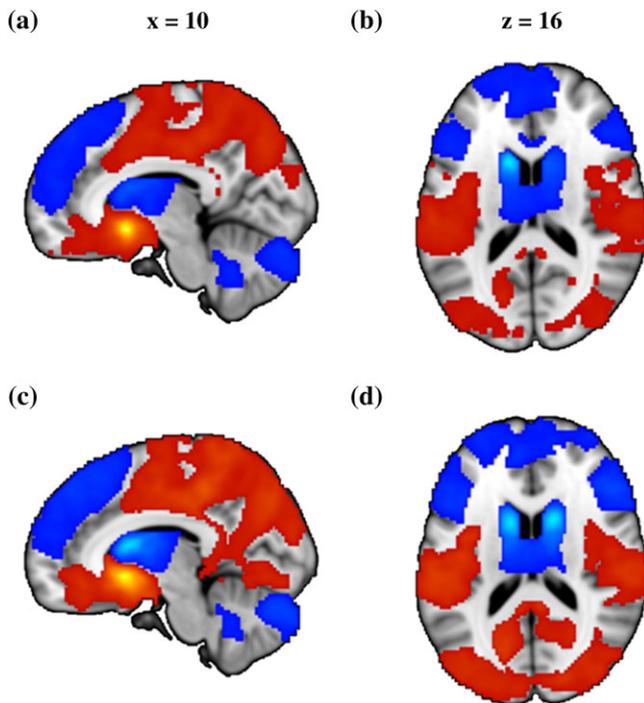
Examining differences in seed-based functional connectivity between the groups revealed no regions showing a significant main effect of group, arguing against unspecific striatal network changes in cannabis dependence. However, significant group  $\times$  seed interaction effects were found mirroring a shift in network level communication of the striatal subregions in cannabis dependence. The interaction effects were observed in connectivity with the dmPFC and rACC across both analyses (Table 3 and Figure 3).

#### 3.4 | Post hoc analyses

To further disentangle the direction of the changes using post hoc comparisons, mean Fisher z-transformed correlation coefficients were extracted from anatomical masks referring to the dmPFC and rACC. Masks of the respective regions were defined using structurally defined regions. To obtain the rACC mask, we first thresholded the anterior cingulate gyrus (HO atlas) at 30% to remove voxels with low probability and the posterior boundary was then delineated as one slice anterior to the coronal plane where the connection of the corpus callosum in each hemisphere was no longer connected (Asami et al., 2008; McCormick et al., 2006). Similarly, to obtain the DMPFC mask, we thresholded the superior frontal gyrus (HO atlas) at 30% probability, excluding voxels greater than 10 mm from the midline of the brain (de la Vega, Chang, Banich, Wager, & Yarkoni, 2016) or posterior to the anterior boundary of supplementary motor area. As displayed in Figure 4a,c, post hoc pairwise comparisons revealed attenuated negative correlation (anticorrelation) between dmPFC and VS ( $t_{[50]} = 3.23$ ,  $p = .002$ , Cohen's  $d = 0.901$ ), as well as decreased positive connectivity between dmPFC and DS ( $t_{[50]} = -2.33$ ,  $p = .024$ , Cohen's  $d = 0.649$ ) in cannabis users relative to controls. Post hoc comparisons



**FIGURE 1** Striatum regions discriminating cannabis-dependent subjects and nonusing controls. Images are thresholded at accuracy  $> 0.75$  (approximate  $p < .001$ , uncorrected) and cluster size  $> 5$  for display purpose



**FIGURE 2** Brain regions showing significant main effect of seed on functional connectivity. (a) and (b) Results from data-driven seeds. (c) and (d) Results from validation seeds. Red color indicated stronger functional connectivity with VS than DS and blue color indicated stronger functional connectivity with DS than VS. All images were thresholded at  $p < .05$ , TFCE corrected

on the rACC demonstrated increased positive functional connectivity with the VS ( $t_{[50]} = 2.32$ ,  $p = .025$ , Cohen's  $d = 0.644$ ) and concomitantly decreased positive functional connectivity with the DS ( $t_{[50]} = -2.69$ ,  $p = .009$ , Cohen's  $d = 0.751$ ) in cannabis users compared to controls.

### 3.5 | Control analyses

Importantly, none of the cannabis users fulfilled a dependence for other illicit drugs, and cannabis was their main drug of abuse. However, in line with an increased prevalence of illicit drug use in addicted populations, the cannabis group had higher experience with

recreational use of other drugs, particularly amphetamine (~60%). Given that previous studies reported altered striatal functioning in occasional amphetamine users (Schrantee et al., 2016), we additionally compared the altered neural indices between users with a history of amphetamine use versus no history. This additional control analysis did not reveal significant between group differences (all  $ps > 0.2$ ), arguing against recreational amphetamine use as driver of the observed difference with the control group. Moreover, the control analysis with the literature-derived validation seeds revealed comparable results as the data-driven seeds, further confirming the robustness of the findings (Figure 4b,d).

### 3.6 | Functional characterization of the altered networks

NeuroSynth decoding revealed that terms showing highest correlations with the dmPFC were predominantly referring to control-related or reasoning-related processes (Figure 5a), while a predominance of reward-related or default network processes (Figure 5b) was found for the rACC.

### 3.7 | Association between cannabis use parameters and neural alterations

Exploring associations between neural alterations and the age of onset of cannabis use, cumulative exposure, and duration of abstinence (days since last use) did not reveal significant associations (all  $ps > 0.1$ ). Corresponding exploratory analyses with nicotine use parameters did not reveal significant associations (all  $ps > 0.2$ ).

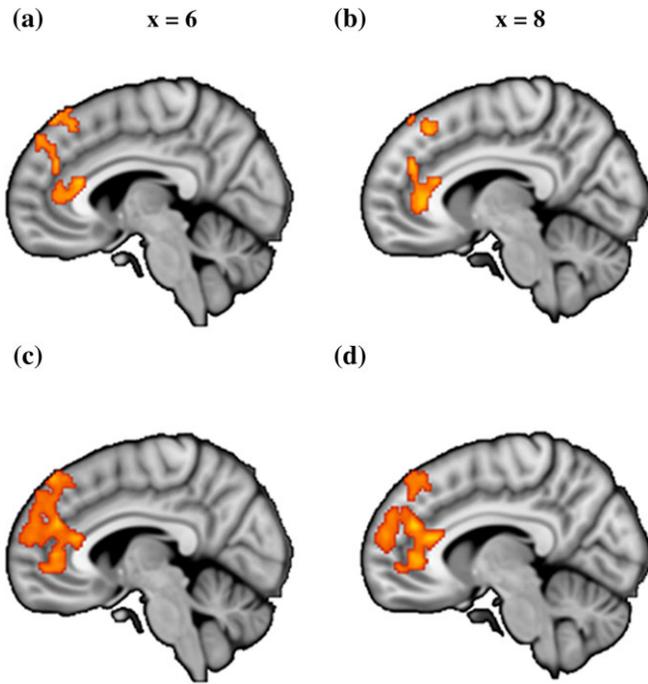
## 4 | DISCUSSION

The present study combined an innovative network level approach with resting state fMRI to determine regional-specific striatal alterations and associated changes in functional communication in abstinent cannabis-dependent subjects. In line with animal and human data that emphasize the specific contributions of the VS and DS, subregion-specific differences between cannabis users and controls mapped to the nucleus accumbens and the caudate nucleus. Cannabis users displayed a relative shift between VS and DS communication

**TABLE 3** Brain regions showing significant group (control vs. user) by seed (VS vs. dorsal striatum) interaction

x	y	z	(sub)Peak voxel F value	p value (TFCE corrected)	k	BA	Laterality	Label
Data-driven seeds								
2	52	42	24.22	0.020	376	9/32/24	L/R	dmPFC
10	36	2	23.40	0.022				rACC
10	30	48	21.42	0.026	163	8	R	dmPFC
6	44	54	16.48	0.039				dmPFC
Validation seeds								
-6	44	20	21.86	0.004	2,147	8/9/10/24/32	L/R	rACC
12	38	2	27.39	0.005				rACC
2	52	42	26.29	0.007				dmPFC

x, y, z = MNI-coordinates; k = cluster size; BA = Brodmann area; dmPFC = dorso-medial prefrontal cortex; DS = dorsal striatum; rACC = rostral anterior cingulate cortex; VS = ventral striatum.



**FIGURE 3** Brain regions showing significant group (control vs user) by seed (ventral vs dorsal striatum) functional connectivity interaction. (a) and (b) Results from data-driven seeds. (c) and (d) Results from validation seeds. All images are thresholded at  $p < .05$ , TFCE corrected

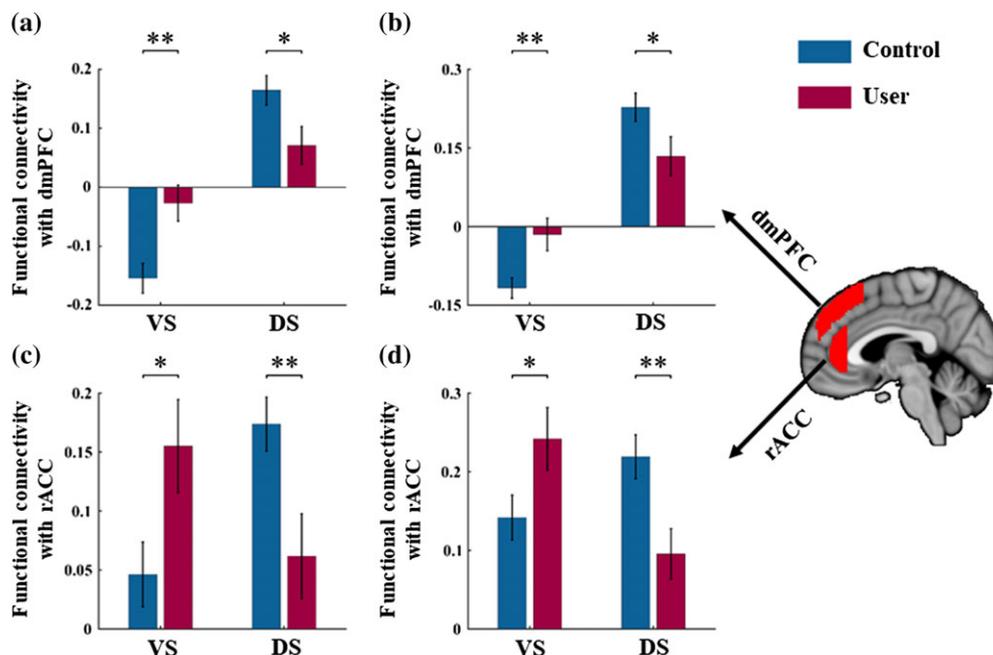
with frontolimbic regions that have been consistently involved in reward processing (rACC) and executive/regulatory functions (dmPFC). Aberrant frontostriatal functional connectivity in the abstinent cannabis users was observed in the absence of a significant main effect in the comparison with the control group. This observation is

partly in line with a previous study reporting that altered frontostriatal connectivity in chronic cannabis users normalizes during 28 days of abstinence (Blanco-Hinojo et al., 2017). However, comparing the shift between the ventral and dorsal striatal functional networks in the present study revealed evidence for a persisting imbalance in the subregion specific frontostriatal circuits. Together, these findings argue against unspecific long-term striatal changes in cannabis dependence but rather emphasize the importance of both, specific adaptations in dorsal and ventral subregions and a relative shift between the regions.

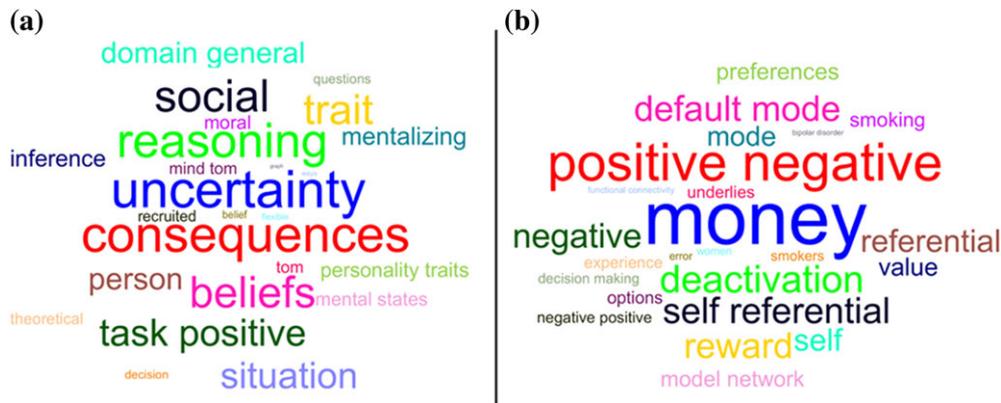
Moreover, employing a fully data-driven network approach that combines the ICC with a pattern classification demonstrated that the general connectivity patterns of the VS and DS can reliably discriminate between cannabis-dependent individuals and controls. As such, the present data are in line with previous studies reporting altered VS and DS activity in chronic cannabis users during drug cue reactivity, reward processing (Weinstein et al., 2016; Wrege et al., 2014; Yanes et al., 2018) and social decision making (Gilman, 2017).

Furthermore, our data emphasize the important contribution of striatal maladaptations in addiction which might reflect a common pathological pathway across addictive disorders. In accordance with a plethora of previous animal and human research suggesting a functional differentiation of the striatum along a dorsal to ventral gradient (Delgado, 2007) and a specific contribution of the striatal subregions to the transition to addiction (Everitt & Robbins, 2016), differences in the intrinsic function of both subregions were demonstrated. Specifically, the nucleus accumbens, a core reward processing node (Delgado, 2007), and the caudate nucleus, strongly engaged in cognitive processes and executive control (Haber, 2016), differed between both groups.

The mapping of the identified regions and their proposed functional contributions was further confirmed by the stronger connectivity of the



**FIGURE 4** Patterns of functional connectivity associated with a significant group (control vs user) by seed (VS vs. DS) interaction. (a) and (c) Results from data-driven seeds and (b) and (d) Results from validation seeds. Error bars reflect the SEM. \* $p < .05$ ; \*\* $p < .01$ ; VS: Ventral striatum; DS: Dorsal striatum; dmPFC: Dorsomedial prefrontal cortex; rACC: Rostral anterior cingulate cortex



**FIGURE 5** Word cloud showing the correlation strength between the ROIs and the meta-analytic map. (a) The top 25 terms associated with dmPFC based on Neurosynth decoding and font size represents relative correlation strength of that term to dmPFC. (b) The same information for rACC

VS with reward-related limbic and (orbito-)frontal regions, whereas the DS showed a stronger functional interaction with frontal regions such as the dorsolateral prefrontal cortex that are critically involved in cognitive and regulatory processes. Examining differences in the connectivity profiles of the identified VS and DS regions revealed a relative shift in the connectivity with the dmPFC, a region that has been implicated in regulatory downstream control over the striatum (Kragel et al., 2018; Robbins et al., 2012) and the rACC, a region engaged in reward processing via up-stream signaling of the VS (Haber & Knutson, 2010). As such, the present findings are in accordance with recent quantitative and qualitative reviews suggesting alterations in networks engaged in cognitive control and reward processing in chronic cannabis users (Weinstein et al., 2016; Wrege et al., 2014; Yanes et al., 2018).

More specifically, both striatal regions demonstrated an uncoupling with dmPFC regulatory regions which may reflect deficient inhibitory control which is consistently observed in cannabis users (Wrege et al., 2014). The altered caudate-dmPFC pathway suggests specific deficits in stop-signal inhibition (Robbins et al., 2012), an inhibitory domain that has been shown to be deficient across addictive disorders (Acikalin, Gorgolewski, & Poldrack, 2017; Morein-Zamir, Jones, Bullmore, Robbins, & Ersche, 2013).

By contrast, in cannabis-dependent subjects the rACC exhibited increased connectivity with the VS and concomitantly decreased connectivity with the DS. Increased connectivity in the nucleus accumbens-ACC pathway in response to drug cues has previously been identified as a specific marker for dependent versus nondependent cannabis users (Filbey & Dunlop, 2014). Activation in the rACC and the adjacent ventromedial PFC has been found to reliably reflect subjective value and reward representations (Acikalin et al., 2017).

The pathway with the VS has been associated with the degree of delayed discounting-associated inhibitory control (Li et al., 2013; see also circuit model by Robbins et al. (2012)) and increased reward-seeking impulsivity in addictive disorders (Fineberg et al., 2010). Consistent with the present results on decreased DS-rACC connectivity, a previous study reported attenuated connectivity in this pathway in nicotine-dependent individuals (Sweitzer et al., 2016). This has been interpreted as disengagement of habitual behavior in favor of conscious perception of craving that promotes drug-seeking behavior.

Together, the pattern of increased connectivity in the VS-rACC pathway engaged in reward-seeking impulsivity and concomitantly decreased striatal coupling with dmPFC regulatory regions may reflect a hyperactive striatal “impulsive” system and a hypoactive prefrontal inhibitory (“reflective”) system as proposed by dual-process models of addiction (Jentsch & Taylor, 1999; Noël et al., 2013). The present findings indicate that the imbalance between the systems extends to the task-free state, suggesting that changes in the intrinsic organization of the striato-frontal circuitries may promote loss of behavioral control in interaction with external stimuli such as drug-associated cues.

Previous animal and human research indicate that alterations in striato-frontal circuits (Becker et al., 2015; Ersche et al., 2012; George & Koob, 2010) and associated functional domains such as deficient regulatory control and increased impulsivity (Dalley, Everitt, & Robbins, 2011; Ersche, Turton, Pradhan, Bullmore, & Robbins, 2010) precede the escalation of drug use and represent predisposing risk factors for addiction. The retrospective design of the present study does not allow to disentangle predisposing alterations from neuroplastic changes that accompany the transition to cannabis dependence. As such, the observed intrinsic striatal alterations may represent predisposing factors that render individuals vulnerable for the development of cannabis dependence.

The present findings need to be considered in the context of limitations. (a) To facilitate the determination of elementary frontostriatal alterations in cannabis dependence the study focused on male participants. The rationale was based on an increasing number of studies reporting modulatory effects of menstrual cycle phase on frontostriatal processing and associated functions (e.g., Diekhof & Ratnayake, 2016; Dreher et al., 2007) as well as recent studies suggesting that addiction-related alterations in these circuits vary as a function of menstrual cycle (Franklin et al., 2015; Wetherill, Jagannathan, et al., 2016; for cannabis see Wiers et al., 2016). The focus on a male sample allowed to control for these potential confounding variables, however, comes at a cost of a limited generalization of the present findings to cannabis-dependent women. In the context of emerging evidence for a higher sensitivity of women to the adverse effects of chronic cannabis use on the brain, including blunted dopaminergic reactivity in the striatum and impaired frontal activity (Paola Castelli et al., 2014; Wiers

et al., 2016), future studies need to consider to explicitly examine sex differences, (b) due to the high prevalence of tobacco use in cannabis addiction both samples included a high proportion of tobacco smokers, although the groups were matched with respect to tobacco use parameters complex interactions between cannabis and tobacco use may have contributed to the present findings, (c) cannabis users that were interested in participation but did not remain abstinent during the 28-day period were not included in the study, the sample thus may represent a subgroup with relative low degrees of cannabis dependence, and (d) the cannabis sample size was relatively small which limited ability to assess within-group clinical correlations.

Together, the present findings suggest a shift in the balance between DS and VS control of behavior in cannabis dependence. These alterations may reflect predisposing risk factors for the development of cannabis dependence or neuroplasticity changes that accompany the transition to dependence. Similar alterations have been previously validated in comprehensive animal models of addiction and may promote the loss of control central to addictive behavior.

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## CONFLICT OF INTERESTS

The authors declare no potential conflict of interests.

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