Neural circuit basis of adolescent THC-induced potentiation of opioid responses later in life

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ABSTRACT:

Use of one drug of abuse typically influences the behavioral response to other drugs, either administered at the same time or a subsequent time point. The nature of the drugs being used, as well as the timing and dosing, also influence how these drugs interact. Here, we tested the effects of adolescent THC exposure on the development of morphine-induced behavioral adaptations following repeated morphine exposure during adulthood. We found that adolescent THC administration potentiated reward, and paradoxically impaired the development of morphine reward. Following a forced abstinence period, we then mapped the whole-brain response to a moderate dose of morphine, and found that adolescent THC administration led to increased morphine-induced activity in the basal ganglia and increased functional connectivity between frontal cortical regions and the ventral tegmental area. Last, we show using rabies virus-based circuit mapping that adolescent THC exposure triggers a long-lasting elevation in connectivity from the frontal cortex regions onto ventral tegmental dopamine cells. Our study adds to the rich literature on the interaction between drugs of abuse, including THC and opioids, and provides potential circuit substates by which adolescent THC exposure influences responses to morphine later in life.

INTRODUCTION:

Cannabis is currently the most widely used drug by adolescents in the United States[1]. Despite a reputation as a relatively harmless drug, over 30% of cannabis users meet the diagnostic criteria for cannabis use disorder (CUD)[2], a similar proportion to those who become addicted after trying "hard" drugs such as cocaine and opioids[3]. With the global trend toward cannabis legalization enabling increased access to cannabis, it is critical to understand how adolescent cannabis use impacts the brain and behavior, especially if it does so in a persistent manner in the dynamically developing adolescent brain. Adolescent exposure to the major psychoactive constituent of cannabis, tetrahydrocannabinol (THC), has been linked to a variety of negative outcomes later in life[4–6], including increased subsequent consumption of 'harder" drugs[7–12]. Given the posited role of the mesolimbic dopamine system as the final common pathway for addiction[13], it is not surprising that brain regions such as the ventral tegmental area (VTA), medial prefrontal cortex (mPFC), and nucleus accumbens (NAc) are critically important in mediating the long-lasting effects of adolescent THC on responses to psychoactive drugs later in life[10,14–16]. Much like use of other psychoactive drugs[17–20], adolescent THC usage can lead to a hyperdopaminergic state[15] that then facilitates a variety of neuroadaptations that promote addiction. Despite this common framework, our mechanistic understanding of how adolescent THC causes persistent changes that linger until adulthood is limited.

Given that THC carries an addictive liability, it is critically important to understand the interactions between adolescent THC usage and exposure to drugs of abuse later in life, for example prescription opioids. It has long been hypothesized that cannabis use may predispose individuals to opioid use[21–23].

This is supported by the observation that cannabinoids and opioids may both activate the midbrain
dopamine system, perhaps via common mechanisms of mu opioid receptor activation[24]. In addition,
cross-sensitization of THC and morphine has been demonstrated[25], adolescent THC exposure can alter
expression of genes related to the endogenous opioid system[10], as well as potentiate opioid-seeking
behaviors later in life[26,27], providing further evidence that these drugs may work via common mechanisms.
This is an important question because, despite the goal to reduce the clinical usage of opioids, opioids
remain the primary treatment for post-operative pain management. In addition, approximately eighty
percent of those who use heroin began using opioids for pain management[28,29]. Therefore, it is important
to understanding mitigate the negative effects that adolescent THC may have on behavioral responses
to opioids later in life.

Here, we administer a moderate dose of THC (5 mg/kg) for 14 days starting at postnatal day 30 (PD30),
during adolescence. After a washout period during which mice develop to adulthood, we then test a
variety of opioid-induced behaviors by administering morphine and assessing differential effects in those
with or without a history of adolescent THC. After undergoing a forced abstinence period following
repeated morphine injections, we also assess how adolescent THC administration influences brain-wide
activity patterns following a moderate of morphine, as well as how adolescent THC administration alters
connectivity onto dopamine neurons in the VTA.

METHODS:
All procedures were approved by the University of California, Irvine's Institutional Animal Care and Use
Committee (IACUC) and carried out in accordance with the National Institutes of Health guidelines for
the care and use of animals. Adolescent male and female C57BL/6J mice were weaned and sexed at
PD21 and group housed with same-sex littermates. Mice were housed in standard individually ventilated
cages with corncob bedding and two cotton nestlets for enrichment. Lights were on a 12 hr on/12 hr off
cycle (7:30-7:30) and rooms have controlled temperature (22°C +/- 2°C) and humidity (55-65%). Mice
received ad libitum access to food and water. Mice were housed in the Gillespie Neuroscience Research
Facility (GNRF), except for during adolescent THC treatment. Mice were transferred to the McGaugh Hall
mouse facility following weaning at approximately P25, and then returned to the GNRF at approximately
P45.

Adolescent THC Administration: Mice were handled and weighed daily. On PD30, mice began daily
injections of either 5 mg/kg THC or vehicle, which was prepared daily by dissolving in 5% Tween80 in
saline. After 14 days of daily injections (PD30-PD43), mice were left in their home cage undisturbed
(aside from weekly cage changes) until PD70 when behavioral testing began.

Drugs: THC was provided by the NIDA Drug Supply Program or Cayman Chemicals (Ann Arbor, MI) and

administered at 5 mg/kg. Morphine was purchased from Patterson and was administered at 10 mg/kg.

Behavioral Testing:

Locomotion: The distance traveled in the open field was assessed during 30-minute recording sessions.

Locomotion was assessed once per day for seven days, the first two following saline injections for

habituation days, and the last five following injection of 10 mg/kg morphine.

Conditioned place preference: Custom two chamber acrylic boxes separated by a small corridor with

distinct visual contexts on the walls of each side were used for CPP tests. On the pre-test day, mice were

placed in the box with no barriers and were allowed to freely explore both sides for 30 minutes, and the

time spent in each chamber was recorded. Following the pre-test, mice underwent two saline pairings

and two morphine pairings. In the morning, the mice were given saline and paired with the preferred

chamber, and in the afternoon, injected with morphine and paired with the non-preferred chamber.

Control mice received saline in both contexts. On the post-test day, mice were again placed in the CPP

chamber with no barriers, and the amount of time spent in each chamber was recorded. Data were plotted

as the relative time spent on the drug-paired chamber in the post-test relative to the pre-test. Mice that

spent over 1200 seconds on one side during the pre-test were excluded from analysis.

Elevated plus maze: Mice were placed at the end of one of the open arms, facing outwards. Mice were

then able to freely explore the maze for 5 minutes, and the time spent in the open arms was recorded.

Tests were performed in dim lighting ~25 lux.

Open field test: Mice were placed in a 40cm x 40cm square arena made of opaque plexiglass and were

able to freely explore during a 5-minute test. During analysis, the arena was divided into nine equal

square zones. The time spent in the center square for each mouse was calculated. Tests were performed

under bright lighting conditions of ~1000 lux.

Marble burying: Mice were placed in a rat cage with 20 marbles placed on top of corn-cob bedding. After

30 minutes, the number of marbles that were buried by each mouse was recorded.

Von Frey: The von Frey test was used to assess mechanical sensitivity. Mice were placed on top of a

wire mesh platform, restrained underneath a glass beaker, and allowed to acclimate for at least 30

minutes. The mouse's right hindpaw was gently touched with different filaments, starting with the 0.4g

filament. If the mouse showed a withdrawal response (such as lifting, shaking, or licking the paw), the

experimenter stepped down to a lighter filament, and if the mouse did not respond, the experimenter

stepped up to a thicker filament until a response occurred. The process continued with filaments of higher

or lower forces for four additional turns following the first change in response. This up/down method was

used to determine the mechanical force required to elicit a paw withdrawal response (50% threshold)[30].

cFos induction and immunohistochemistry: One hour following injection of 5 mg/kg morphine (a dose

used only for morphine-induced reinstatement), mice were perfused with PBS followed by 4%

formaldehyde, and brains isolated and post-fixed overnight in 4% formaldehyde. The following day, brain

clearing began using iDISCO[31]. The monoclonal rat anti-cFos antibody (Synaptic Systems lot #226017)

in 5% NDS/PBS-Tween 0.1% was used at a 1:5000 dilution, with an incubation time of five days. Donkey

anti-rat Cy3 (Jackson Immunoresearch) was then used at a dilution of 1:250. Brains were then cleared

in dibenzyl ether and imaged using a Zeiss Z1 light-sheet microscope. ClearMap[32] was then used to align

the imaged brains to the Allen Brain Reference Atlas, and to quantify the number of cFos+ neurons in

anatomically-defined brain regions.

Network Analysis: In order to relate functional connectivity using cFos labeling to anatomical connectivity,

anatomical projection data from the Allen Mouse Brain Connectivity Atlas was used to construct a

connectivity matrix among the brain regions in our analysis[33]. This atlas quantifies axonal arborization

from AAV injections to measure projections from one region to another. The density of each projection,

or the sum or projecting pixels divided by the total pixels in that region, was normalized by dividing by the

volume of injection, so as not to bias experiments with more virus injection than others. This quantity was

also multiplied by the number of voxels in the injection structure to avoid overrepresentation of smaller

regions. This quantity was computed for every pair of source and target regions to build a connectivity

matrix. We log-normalized these quantities so that they would have a more normal distribution and

behave better in network analysis. Community detection was performed using the Louvain algorithm with

a resolution parameter of 1, in the scit-kit network library[34]. Once the clusters were determined, the cFos

fold change of their regions were compared using a one-way ANOVA followed by pairwise t-tests.

Functional Analysis: Regional cFos counts across each mouse in the adolescent vehicle-treated and

THC-treated conditions were dimensionally reduced using UMAP, so that regions with similar counts

across the mice would be embedded close to each other[35]. Regions were clustered using HDBSCAN,

which can identify hierarchical clusters in continuous data based on variations in density of the data[36].

Hierarchical clustering was also performed on the correlations between regions, to look more closely at

their functional relationships. R^2 value from Pearson correlations were computed for each pair of regions,

based on their counts across mice of a given condition. Hierarchical clusterings were computed on

correlation matrices using the average linkages methodology. For visualizations, the hierarchical

clustering tree was cut at a height of 0.75 to determine the clusters (Figure 4I-J). The trees were cut for

heights between 0.5 and 1 to observe trends in modularity across the two conditions. The seaborn library
was used for all hierarchical clustering computations[37].

Rabies virus circuit mapping: Rabies virus (RABV) monosynaptic tracing studies were carried out as
previously described[19,38], with minor modifications. On Day 0, 100 nL of a 1:1 volume mixture of AAV-
CAG-FLEx^{loxP}-TC and AAV-CAG-FLEx^{loxP}-G was injected into the VTA of PD56 mice that had undergone
two weeks of THC injections between PD30 and PD43. Fourteen days later, RABV was injected into the
VTA. After recovery, mice were housed in a BSL2 facility for 5 days to allow RABV spread and GFP
expression. Mice were then perfused with PBS followed by 4% formaldehyde and postfixed overnight.
Fixed brains were cryoprotected by incubation in a 30% sucrose solution overnight, and frozen in OCT.
Brains were then serially sliced on a cryostat into 60 um sections. GFP+ cells throughout the brain in 22
anatomically-defined input sites were counted manually, as previously performed[19,38].

Viruses and injection coordinates: Coordinates used for the VTA were (relative to Bregma, midline, or
dorsal brain surface and in mm): AP − 3.20, ML 0.4, DV − 4.2.

The titers of viruses, based on quantitative PCR analysis, were as follows:
AAV-CAG-FLEx^{loxP}-TC, serotype 5, 2.4 x 10^{13} genome copies (gc)/ml
AAV-CAG-FLEx^{loxP}-G, serotype 8, 1.0 x 10^{12} gc/ml

For RABV, the titer was performed on HEK 293T-TVA800 cells.
RABV (EnvA-pseudotyped, GFP-expressing), 5.0 x 10^{8} colony forming units (cfu)/ml

After scaling RABV data, PCA was applied so features were brain regions and observations were
individual brains of mice treated with various drugs. Differences between mouse brains were
visualized by plotting PC1, PC2, and PC3 (Figure 5J-L, Q-R). Ellipsoids for each condition were
calculated and overlayed onto plots to visualize trends in PC values by condition. To further quantify
relationships between brains and conditions, the Euclidean distance between each point (or each
brain) in the PC1 versus PC2 coordinate space was calculated. These distances were then plotted
in a heatmap with brains being organized by similarity assessed by hierarchical clustering (Figure
5M). Alternatively, for a summary by condition, these distances were averaged based on drug
administered, and the resulting heatmap and clustering were made based on averaged Euclidean
distance values (Figure 5S). Heatmaps and clustering were made using the clustermap function in
the Python package seaborn with default parameters[37]. The RABV data that included single
injections of psychostimulants, morphine, or nicotine, or two anesthetic doses of ketamine/xylazine,

were presented in our published work[39] and used here for sake of comparison. For these
experiments, a single dose of 15 mg/kg cocaine, 1 mg/kg amphetamine, 2 mg/kg methamphetamine,
0.5 mg/kg nicotine, or 10 mg/kg morphine were given to adult mice, and RABV tracing was performed
the following day. These mice were anesthetized with isoflurane-based anesthesia, as used in this
manuscript. A separate group was anesthetized with a ketamine/xylazine mixture prior to both helper
virus injection and RABV injection, and not given another drug. RABV was introduced during the
second ketamine/xylazine injection.
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Quantification and Statistical Analysis: Statistics for all studies were calculated using GraphPad Prism
9 software. Statistical significance between direct comparisons was assessed by unpaired or paired t-
tests. When multiple conditions were compared, one- or two-way ANOVAs were first performed, as
appropriate, then t-tests were then performed for each individual comparison. Multiple comparisons
corrections were performed when multiple such t-tests were being performed e.g., RABV input mapping
and cFos quantification, and significance was assessed using the Holm-Bonferroni method. In conditions
where multiple comparisons were performed and the results were still considered significant, asterisks
were presented corresponding to the original p-values. Paired t-tests were used in cases with repeat
measurements. Dot plots presented throughout the manuscript include a bar representing the mean value
for each group. Error bars represent s.e.m. throughout. For all figures, ns $P > 0.05$, * $P \leq 0.05$, ** $P \leq 0.01$,
*** $P \leq 0.001$, **** $P \leq 0.0001$.
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RESULTS:
Adolescent THC exposure history reduces morphine-induced locomotion, with no impact on
morphine reward
We first exposed mice to a daily 5 mg/kg/day dose of THC (or its vehicle) during adolescence, here
defined as postnatal days 30-43 (Figure 1A). This is considered a moderate dose that is comparable to
doses self-administered by humans[40,41]. Following vehicle or THC administration, mice were left in their
home cages undisturbed besides weekly cage changes until postnatal day 70, at which time behavioral
testing commenced. The protocol of morphine administration and behavioral testing is shown in Figure
1B. To test how adolescent THC exposure history impacts morphine reward, we conducted a conditioned
place preference (CPP) test using 10 mg/kg morphine (Figure 1C). We induced a significant preference
to the morphine-paired side in both vehicle and THC exposed animals, to an approximately equivalent
extent. No preference was observed in animals treated with vehicle during adolescence and again at
adulthood in place of morphine (Vehicle/saline 762s pre, 778s post, $p = 0.22$; Vehicle/morphine 783s pre
vs. 887s post, $p = 0.038$; THC/morphine 785s pre vs. 887s post, $p = 0.0017$; Figure 1D). To determine
whether adolescent THC treatment impacted morphine-induced locomotion, we measured locomotor

behavior while mice were in an open field. Following a two-day habituation protocol where mice were

given saline injections, 10 mg/kg morphine was given once per day for five days (Figure 1E) to test the

sensitization or tolerance to morphine's effects on locomotion. We observed that morphine increased

locomotion the most following the first dose, and this elevation was reduced with each subsequent dose,

for both groups. We also found that that adolescent THC exposure overall blunted morphine-induced

locomotion during the five-day morphine administration window (repeated measures Two-way ANOVA p

= 0.045; Figure 1F), though no individual day was significantly different between the groups following

multiple comparisons corrections. These data indicate that adolescent THC exposure reduced morphine-

induced locomotion following morphine exposure later in life.

Adolescent THC exposure impairs the development of anxiety-related behavior and pain

hypersensitivity during morphine withdrawal.

We next assessed whether mice treated with THC during adolescence were differentially impacted by

withdrawal from repeated morphine administration. Following the five days of morphine given for

locomotor testing and two days for CPP extinction, mice were left in their home cages for an additional 8

days, for a total abstinence period of ten days (Figure 2A). Following this ten-day forced abstinence, mice

were tested for anxiety-like behavior using three tests: the elevated plus maze, open field test, and marble

burying tests. The elevated plus maze tests how mice balance their tendency to explore against their

preference for enclosed spaces (Figure 2B). A reduction in the time spent exploring the open arms is

interpreted as anxiety-like behavior. Mice treated with repeated morphine injections spent less time in

the open arms of the elevated plus maze relative to those treated with repeated saline injections. However,

adolescent THC exposure blocked the reduction in time spent in the open arms induced by repeat

morphine injection (One-way ANOVA p = 0.0005; pairwise t-tests, vehicle/saline vs. vehicle/morphine

47.73s vs. 17.84s, p = 0.01; vehicle/saline vs. THC/morphine 47.73s vs. 42.39s, p = 0.67;

vehicle/morphine vs. THC/morphine 17.84s vs. 42.39s, p = 0.0018; Figure 2C). This suggests that

adolescent THC exposure suppressed withdrawal-induced anxiety behaviors in response to adult opioid

exposures. Similar results were observed using the open field test, where mice are both compelled to

explore new environments, and avoid open areas (Figure 2D). Less time spent exploring the center of

the arena is interpreted as anxiety-like behavior. We found that repeat morphine injections reduced time

spent in the center of the open field, and THC history reduced this anxiety-like behavior (One-way ANOVA

p = 0.027; pairwise t-tests, vehicle/saline vs. vehicle/morphine 22.66s vs. 12.55s, p = 0.030;

vehicle/saline vs. THC/morphine 22.66s vs. 15.85s, p = 0.077; vehicle/morphine vs. THC/morphine

12.55s vs. 15.85s, p = 0.59; Figure 2E). Finally, in the marble burying test mice tend to bury marbles that

are exposed in the bedding of a new cage (Figure 2F). In this assay, more marbles buried is interpreted

to reflect increased anxiety-like behavior. We found that mice undergoing forced abstinence following

repeat morphine injections buried more marbles after the 30-minute test, whereas adolescent THC

administration reduced this number, consistent with a reduced anxiety-like behavior (One-way ANOVA p
= 0.0006; pairwise t-tests, vehicle/saline vs. vehicle/morphine 0.93 vs. 3.67, p = 0.0048; vehicle/saline
vs. THC/morphine 0.93 vs. 0.50, p = 0.83; vehicle/morphine vs. THC/morphine 3.67 vs. 0.50, p = 0.006;
Figure 2G). Therefore, the results from each of these tests support the conclusion that adolescent THC
exposure decreases the expression of anxiety-like behaviors in morphine-treated mice.

Next, we used a von Frey assay to examine the effects of adolescent THC exposure on mechanical
hypersensitivity, both before and after morphine injection (Figure 2H). In addition to an increase in
anxiety-like behavior, mice undergoing physiological withdrawal following repeated opioid administration
show an increased pain sensitivity[42,43]. Mice treated with repeated THC doses during adolescence
showed no difference in mechanical thresholds when tested one day prior to morphine CPP testing
(Vehicle/morphine vs. THC/morphine 0.85g vs 0.86g, p = 0.93; Figure 2I). However, when tested again
following repeat morphine exposure and forced abstinence, mice given repeated morphine administration
showed an increased mechanical hypersensitivity relative to saline-treated controls, and adolescent THC
treated reduced the extent of this hypersensitivity (One-way ANOVA p = 0.0032; pairwise t-tests,
vehicle/saline vs. vehicle/morphine 1.09g vs. 0.42g, p = 0.0027; vehicle/saline vs. THC/morphine 1.09g
vs. 0.69g, p = 0.052; vehicle/morphine vs. THC/morphine 0.42g vs. 0.69g, p = 0.30; Figure 2J). Together,
these results indicate that adolescent THC exposure reduces the expression of morphine withdrawal-
induced behaviors following repeated morphine administration during adulthood.

Adolescent THC exposure enhances drug-induced reinstatement of morphine CPP
Following forced abstinence, we next assessed whether mice differentially reinstated CPP following a 5
mg/kg exposure of morphine in the CPP box. Drug exposure following abstinence can reinstate
previously extinguished drug-seeking behaviors, and therefore can serve as a model of relapse in mice.
Both groups of morphine-treated mice showed normal CPP (e.g., Figure 1D). Following CPP, mice then
underwent extinction. To reduce confounding factors while examining reinstatement, mice were excluded
that did not show CPP and proper extinction (N = 16 vehicle, 11 THC; Figure 3A-B). While mice treated
with vehicle during adolescence did not significantly reinstate CPP following the priming 5 mg/kg
morphine injection using this protocol, mice treated with THC during adolescence did significantly
reinstate their CPP (Vehicle/morphine-treated mice 722s vs 801.3s, = p = 0.43; THC/morphine treated
mice 780.7s vs. 1015s, p = 0.027; Figure 3C). This means that adolescent THC exposure drives
reinstatement of drug-seeking behavior in response to a sub-threshold dose of morphine in mice given
vehicle during adolescence.

Whole-brain analysis shows differences in brain activity following drug-primed
reinstatement in adolescent vehicle- vs. THC-treated mice

Given that adolescent THC-treated mice showed morphine-induced reinstatement whereas vehicle-
treated controls did not, we next assessed what neural pathways may be responsible for this effect. To
do this, we perfused the mice 60 minutes following the reinstatement test, cleared the brains using
iDISCO+, and immunostained for the immediately early gene cFos to obtain a brain-wide map of neuronal
activity (Figure 4A). We observed an overall elevation of neuronal activity in response to 5 mg/kg
morphine in mice treated with THC during adolescence than vehicle, as evidenced by a higher number
of cFos+ cells upon morphine exposure (Figure 4B). This increase was exhibited in most brain regions,
with some of the highest-fold elevations occurring in basal ganglia regions including the globus pallidus
externus (GPe), subthalamic nucleus (STN) and substantia nigra pars reticulata (SNr), as well as areas
such as the substantia innominate/ventral pallidum, and several thalamic regions (Figure 4B-D). In fact,
only four brain regions showed a reduction in cFos labeling in adolescent THC-treated mice, which
included both the medial and lateral habenula.

As the elevation in cFos labeling occurred globally in the mouse brain, we wanted to assess whether this
elevation was larger for some pathways or whether the activation was uniform. To test this, we built a
quantitative connectivity matrix of the mouse brain, using the Allen Mouse Brain Connectivity Atlas as a
template[33]. Doing so revealed the existence of four major network communities within the brain, largely
consisting of subcortical regions including the VTA (community 1, purple), cortical regions (community 2,
blue), the hippocampus and related connections (community 3, red), and mid/hindbrain regions
(community 4, yellow) (Figure 4E). We then assessed the relative increase in cFos labeling of regions in
each community in adolescent THC- vs. vehicle-treated mice following a reinstatement opioid dose. We
found a significant increase in the log fold-change cFos labeling in the purple and blue communities
relative to the yellow community, suggesting that the subcortical and cortical modules both showed an
overall larger increase in activity relative to mid/hindbrain regions in THC-treated mice (Figure 4F).

Following this analysis, we wanted to assess if these community-specific changes in cFos activation were
accompanied by any changes in the functional relationships of different brain regions. To do so, we took
the cFos labeling data across all regions for all mice and performed a dimensionality reduction analysis
using Uniform Manifold Approximation and Projection (UMAP). UMAP divided the brain regions into
approximately 10 activity-defined clusters in the adolescent THC-treated mice data. These clusters which
mapped relatively consistently onto the adolescent vehicle-treated mice data (Figure 4G-H), with a larger
distribution of brains from vehicle-treated compared to THC-treated mice. Interestingly, while frontal
cortex regions, including the mPFC subregions infralimbic cortex (IL), prelimbic cortex (PL), anterior
cingulate cortex (ACC), and anterior insula cortex (AI) all were in relatively close proximity in UMAP space
in both adolescent vehicle-treated mice and THC-treated mice, they clustered much more tightly together
in THC-treated group relative to the vehicle-treated group, indicating a higher level of correlated activity

following a reinstatement opioid exposure (Figure 4G-H). This suggests that adolescent THC exposure

may increase the functional connectivity between frontal cortex regions and the VTA. To assess

functional modularity more quantitatively, we built a brain-wide functional correlation matrix for adolescent

THC-treated and vehicle-treated mice based on their cFos labeling (Figure 4I-J). While in vehicle-treated

mice the VTA and frontal cortex regions separated into different clusters, reflecting their separation into

different networks (Figure 4I), in THC-treated mice the IL, PL, ACC, and AI all co-clustered with the VTA,

indicating a higher level of correlated activity amongst these regions in adolescent THC-treated mice

(Figure 4J). Last, we wanted to assess the overall modularity of brain activity in adolescent THC- vs.

vehicle-treated mice. To do this, we determined the number of unique clusters across a range of

parameters. Across a wide parameter range, adolescent THC-treated mice exhibited fewer activity-

defined clusters than vehicle-treated mice, reflecting an overall reduced modularity in brains from

adolescent THC-treated mice (Figure 4K). These findings overall show that the increased activation of

cortical and subcortical pathways in adolescent THC-treated vs vehicle-treated mice after reinstatement

is accompanied by a collapse in modularity of pathways in adolescent THC-treated mice, including a

tighter clustering of the VTA with frontal cortex regions.

Adolescent THC exposure changes connectivity to ventral tegmental area dopamine cells

VTA is a critical brain region for the development of a variety of drug-induced behavioral adaptations,

including drug reward, sensitization, and withdrawal. To test whether adolescent THC exposure modified

the input control to VTA dopamine (VTA^{DA}) cells, we mapped inputs to these cells using the rabies virus

(RABV) monosynaptic input mapping method. First, in animals treated with THC during adolescence,

Cre-dependent adeno-associated viruses (AAVs) expressing TVA and the RABV glycoprotein, RABV-G,

were injected into the VTA of DAT-Cre mice. Two weeks later, mice were injected with a glycoprotein-

deleted, EnvA-pseudotyped, GFP-expressing RABV (Figure 5A). This virus infected TVA-expressing

cells by nature of the EnvA-TVA interaction. Once in VTA^{DA} cells, this virus could spread to their inputs,

as RABV-G was expressed in DAT-Cre cells. However, once in input cells, since RABV did not express

its own glycoprotein, the virus could not spread further. Five days later, experiments were terminated,

and GFP-positive cells were counted throughout the brain as we did previously[19,39], and input connectivity

was assessed. RABV-labeled input cells were observed throughout the brain, as in previous studies

(Figure 5B-C)[19,39]. Overall, we observed several quantitative differences, shown in Figure 5D (Two-way

ANOVA, interaction p = 0.0062). While several brain regions showed a visible difference in connectivity,

for example elevated connectivity from cortical regions comprising the anterior cortex (infralimbic cortex,

prelimbic cortex, anterior insula cortex, anterior cingulate cortex, and the anterior motor cortex[19,38,39,44]),

parabrachial nucleus (PBN) and deep cerebellar nuclei (DCN), only the anterior cortex was significantly

different following correction for multiple comparisons (corrected p-value p = 0.029).

This traditional approach to assessing differences in connectivity is statistically conservative, and only

assesses potential differences in connectivity, one input at a time. To reveal further patterns in the data,

we have recently employed a dimensionality reduction approach that enables us to detect larger-scale

differences in input patterns between brains of mice undergoing a variety of treatments[39,45]. We therefore

first used this approach to assess which inputs best differentiated THC-treated vs. control mice using

principal component analysis (PCA). We found that 3 PCs were sufficient to explain approximately 75%

of the variance in the data, and thus we focused only on these 3 PCs (Figure 5E-F). We found that PC1

completely separated THC-treated and control mice, whereas PC2 and PC3 did not (Figure 5G-I). Control

mice had high PC1 values, which were driven predominately by strong contributions from nucleus

accumbens medial shell (NAcMed), core (NAcCore), ventral pallidum (VP), septum, lateral habenula

(LHb) and medial habenula (MHb), while THC-treated mice had low PC1 values, which were driven by

the bed nucleus of the stria terminalis (BNST), extended amygdala area (EAM), entopeduncular nucleus

(EP), paraventricular nucleus of the hypothalamus (PVH), lateral hypothalamus (LH), and PBN (Figure

5G). To quantify these differences, we plotted each brain's location on each PC, and performed t-tests.

Differences between THC-treated and control groups were strongly significant along PC1, and not

different along PCs 2 and 3, as expected (Control 3.3 vs. THC -2.2, p = 0.0002). To further quantify

relationships between brains and conditions, the Euclidean distance between each brain in the PC1

versus PC2 coordinate space was calculated, and the relationships plotted as a correlogram. THC-

treated and control brains separated largely by condition, as expected (Figure 5M).

Last, we assessed how VTA^{DA} connectivity of THC-treated mice compares to mice treated with one of

several drugs, including a single dose of the psychoactive but non-addictive fluoxetine, two doses of the

anesthetic mixture of 100 mg/kg ketamine and 10 mg/kg xylazine, spaced 2 weeks apart, or one dose of

the addictive cocaine, amphetamine, methamphetamine, morphine, or nicotine, together grouped for

simplicity as "drugs of abuse". Using a similar PCA-based assessment, we observed that mice treated

with a drug of abuse separated from the other conditions along PC2, but not PC1 or PC3 (Figure 5N-P).

Interestingly, THC-treated mice had even higher values of PC2 than mice treated with one of the drugs

of abuse and were thus located further along PC2 from controls (Figure 5O-P). Positive values of PC2

were largely driven by strong contributions from the anterior cortex, BNST, EP, CeA, LH, PBN, and DCN,

notably many of the same regions that differentiated brains of THC-treated from controls (Figure 5G, N).

While PC3 did not strongly differentiate most of the groups, it did differentiate THC-treated mice from

those treated with a drug of abuse (Figure 5R). THC-treated mice had higher PC3 values, which were

driven by strong contributions from the anterior cortex, dorsal striatum (DStr), PVH, CeA, PBN, and DCN

(Figure 5N). The locations of each brain in PC space were plotted on bar graphs, and differences were

again assessed (PC2; Saline vs. THC -2.43 vs. 2.96, p < 0.0001; Fluoxetine vs. THC -1.55 vs. 2.96, p =

0.0004; K/X vs. THC -1.23 vs. 2.96 p = 0.0024; Drugs vs. THC 0.19 vs. 2.96, p = 0.0051. PC3; One-way

ANOVA p = 0.0004; Saline vs. THC 0.70 vs. 1.85, p = 0.63; Fluoxetine vs. THC 0.88 vs. 1.85, p = 0.71;

K/X vs. THC 0.19 vs. 1.85 p = 0.27; Drugs vs. THC -0.79 vs. 1.85, p = 0.0005; Figure 5Q-R). The distinct

connectivity patterns were also apparent following our Euclidean distance analysis in PC1 and PC2 space,

which indicated that the connectivity patterns from brains from THC-treated mice segregate completely

from those of mice treated with other psychoactive drugs (Figure 5S). Together, these results indicate

that adolescent THC treatment induces similar but distinct input changes onto VTA^{DA} neurons as previous

exposure to a single exposure to a "hard drug", principally an increase in connectivity from the anterior

cortex.

DISCUSSION:

Here, we tested the effects of adolescent THC exposure on later-life morphine-induced behaviors, brain-

wide activity following morphine reinstatement, and midbrain dopamine cell synaptic connectivity. We

found that adolescent THC exposure blunts morphine-induced locomotion and withdrawal anxiety/pain

hypersensitivity and enhances morphine-induced reinstatement following a forced abstinence period.

This broad range of effects is consistent with findings from studies that have explored the influence of

THC on morphine-induced behavioral changes[14,25–27,46]. Here, we extend these previous studies by

linking brain-wide activity patterns following a reinstatement dose of morphine to morphine-induced

reinstatement behavior and show that adolescent THC exposure creates long-lasting changes in input

connectivity to VTA^{DA} cells. Our study thus adds to a rich literature on the subject by implicating several

discrete circuits as potential drivers of the long-lasting effects of adolescent THC exposure on behavioral

and brain responses to opioids later in life.

It is perhaps surprising that THC exposure blunts morphine withdrawal-induced anxiety across several

behavioral assays, as well as pain sensitivity. Interestingly, the behavioral profile is consistent with the

behavioral responses observed following exposure of drugs of abuse during adolescence[47–51], where

adolescent mice experience potentiated drug reward with only mild withdrawal symptoms relative to adult

mice receiving equivalent drug exposures. This effect could be mediated by an occlusion of opioid-

induced epigenetic changes by those elicited following adolescent THC exposure. Such an effect was

previously observed following juvenile exposure to oxycodone; in this case, oxycodone induced long-

lasting epigenetic changes, for example an elevation of the repressive histone modification H3K27me3

on genes that control dopamine transmission in the VTA[52]. Therefore, one hypothesis is that adolescent

THC exposure may result in a similar imbalance of reward/aversion as is seen in responses to drugs

during adolescence by occluding compensatory responses to drug exposure that typically occur in

adulthood, but not adolescence. This could result, in, for example, withdrawal from repeated opioid

injection failing to induce a compensatory induction in anxiety-related behaviors and pain hypersensitivity.

In this case, reversal of these epigenetic marks may restore mice to basal levels of susceptibility. Future

studies are required to identify the potential cellular loci of these effects, as well as what the epigenetic
consequences of early life THC exposure may be.

In addition, we found that adolescent THC administration induces higher levels of drug-induced
reinstatement, consistent with previous observations assessing drug- or stress-induced
reinstatement[8,9,26]. Here, we extend these observations by linking these results to brain-wide cFos
activation in adolescent vehicle- vs. THC-treated mice. Overall, we found that adolescent THC-treated
mice had higher numbers of cFos+ cells in response to a 5 mg/kg reinstatement dose of morphine than
adolescent vehicle-treated controls. In addition, we found that adolescent THC-treated mice have higher
levels of correlated activity in mPFC and AI regions with the VTA than adolescent vehicle-treated mice,
indicating enhanced functional connectivity between these regions (Figure 4G-H). We previously showed
that activation of these cortical inputs to the VTA promotes reinforcement through increasing dopamine
release in the NAc[38]. Therefore, the elevated functional connectivity that we observe between these
regions in adolescent THC-treated mice may promote greater levels of dopamine release in the NAc than
in adolescent vehicle-treated mice and is one mechanism by which adolescent THC-treated mice may
more strongly reinstate morphine CPP following a priming dose of morphine.

However, one limitation of cFos-based analyses is that they do not consider anatomical connections.
Viral methods such as the monosynaptic RABV circuit mapping technique can specifically label
synaptically-connected cells[53,54]. Furthermore, we recently showed that the extent of input labeling is
dependent on the activity of input connections[19,20,55]. Therefore, our RABV mapping experiments have
the unique ability to explore the changes in connectivity to VTA^{DA} cells following repeated THC exposure,
without a preconceived bias as to which inputs may be altered. We have previously used this assay to
define how drugs of abuse change inputs to VTA^{DA} cells, enabling us to identify several pathway-specific
changes that have important consequences for the development of drug-induced behavioral
adaptations[39]. Here, we compare connectivity of VTA^{DA} cells in brains from mice receiving our adolescent
THC protocol to those receiving a single drug of abuse. We found that adolescent THC exposure induced
numerous long-lasting input changes onto VTA^{DA} cells that partially, but not completely, overlapped with
those induced by "hard drugs". The most prominent difference between mice given adolescent THC
exposure vs. a drug of abuse is that THC-treated mice show a pronounced increase in inputs from the
anterior cortex (Figure 5D). Heightened activation of this connection is consistent with our cFos results
and would be expected to be rewarding through elevating activity in VTA^{DA} cells. This hypothesis is
consistent with the hyper-dopaminergic state of THC-treated animals previously reported[15], and the more
robust response in these animals to a reinstatement dose of morphine. In addition, we observed an
overall reduced modularity in adolescent THC-mice relative to adolescent vehicle-treated controls (Figure
4K). A reduction in brain modularity is seen following use and withdrawal from drugs of abuse[56–61] as well

as brain disorders such as traumatic brain injury and Alzheimer's-related dementia[62–67]. How this reduced
modularity may influence behavioral responses to drugs, however, remains largely unknown.

In our study, we also tested the hypothesis that adolescent THC exposure alters connectivity onto VTA
dopamine (VTA^{DA}) cells. This hypothesis was supported by the observation that rats treated with THC
during adolescence develop a hyper-dopaminergic state[15]; one potential mechanism that could mediate
this effect would be a change in connectivity to facilitate an increase in DA release in downstream brain
regions. Here, we found an elevation in inputs from the anterior cortex to VTA^{DA} cells, which we previously
showed promotes reinstatement by facilitating DA release in the NAc. Notably, we previously have shown
that a single injection of a variety of drugs of abuse can cause long-lasting changes in input connectivity
onto VTA^{DA} cells[19,20,55], with this signature being relatively similar regardless of the drug administered[19,39],
suggesting that these drugs induce similar changes in connectivity. The anterior cortex, particularly the
mPFC, is thought to play important roles in the later stages of substance use disorder[68–71]. The reduction
in connectivity from the anterior following a single dose, and increase in connectivity following repeated
doses, is thus consistent with that hypothesis of altered mPFC engagement during the addiction cycle.
One additional point of note is that we previously showed that drugs of abuse increase connectivity from
the globus pallidus externus (GPe) to the ventral midbrain, and that these changes are important for drug-
induced behavioral adaptations including reward and sensitization[19,20]. However, THC treatment did not
induce a notable change in GPe inputs onto VTA^{DA} cells (Figure 5D). Thus, while THC triggers several
long-lasting changes in connectivity to VTA^{DA} cells, these appear to be distinct from those initiated by
psychostimulants, nicotine, or morphine. However, the conclusions that we can draw from these
experiments are limited by differences in experimental design. For example, for the drug-treated mice,
RABV was injected one day following injection of the drug, whereas in THC-treated mice, they were
injected approximately 35 days following completion of a 2-week daily injection protocol. Thus, while in
the former study design drug-induced changes do not need to persist for longer than a few days to yield
the observed changes, in THC-treated animals the effects observed had to persist for longer than 35
days. In addition, THC was given fourteen times, once per day, whereas all other drugs were only given
once in total. Therefore, the effects observed with THC may result from repeated injections. However,
we performed the experiments presented here because our behavioral assessments were done 35 days
following completion of THC administration, and we wanted the RABV experiments to follow the same
timeline. Further experiments will be needed to resolve these questions.

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FIGURES:

Figure 1: Effects of adolescent THC exposure on behavioral responses to morphine
administration later in life.

A, Timeline of THC or vehicle injections during adolescence. Injections were repeated once per day for
14 total days.

B, Morphine injections, and behavioral testing. The entire protocol is shown for each of Figures 1-3 for
reference, with the relevant tests presented in the figure in full opacity and surrounded by red boxes.

C, Schematic of the conditioned place preference test.

D, Time spent in the morphine-paired chamber in the pre-test and post-test (in the vehicle/saline-paired
group, both chambers were paired with saline). Vehicle/saline 762s pre, 778s post, p = 0.22;
Vehicle/morphine 783s pre vs. 887s post, p = 0.038; THC/morphine 785s pre vs. 887s post, p = 0.0017.

E, Schematic of the locomotor test.

F, Mice treated with THC during adolescence showed an overall reduction in morphine-induced
locomotion relative to mice treated with saline during adolescence. Repeat measures Two-way ANOVA
between conditions $p = 0.045$; Sidak's multiple comparisons tests morphine day 1 $p = 0.68$, morphine
day 2 $p = 0.24$, morphine day 3 $p = 0.31$, morphine day 4 $p = 0.50$, morphine day 5 $p = 0.75$.
Vehicle/saline-treated mice are shown for comparison but were not included in statistical tests. Two-way
ANOVA only included morphine injection days, not habituation days. For this and all figures, error bars =
SEM, ns $p > 0.05$, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$.

Figure 2: Adolescent THC administration reduces expression of morphine-induced withdrawal behaviors during a prolonged abstinence period.

A, Timeline of experiments.

B, Schematic of the elevated plus maze test.

C, Repeated morphine administration leads to a reduction in the time spent in the open arms of an elevated plus maze, indicative of anxiety-like behavior relative to mice treated with saline during adolescence, and saline during adulthood. Adolescent THC exposure prevented this morphine-induced anxiety. One-way ANOVA p = 0.0005; pairwise t-tests, vehicle/saline vs. vehicle/morphine 47.73 s vs. 17.84 s, p = 0.01; vehicle/saline vs. THC/morphine 47.73 s vs. 42.39 s, p = 0.67; vehicle/morphine vs. THC/morphine 17.84 s vs. 42.39 s, p = 0.0018.

D, Schematic of the open field test.

E, Repeated morphine administration leads to a reduction in the time spent in the center of an open field
relative to mice treated with saline during adolescence, and saline during adulthood. Adolescent THC
exposure blunted this reduction. One-way ANOVA p = 0.027; pairwise t-tests, vehicle/saline vs.
vehicle/morphine 22.66 s vs. 12.55 s, p = 0.030; vehicle/saline vs. THC/morphine 22.66 s vs. 15.85 s, p
= 0.077; vehicle/morphine vs. THC/morphine 12.55 s vs. 15.85 s, p = 0.59.
F, Schematic of the marble burying task.
G, Repeated morphine administration leads to an elevation in the number of marbles buried, indicative
of anxiety-like behavior relative to mice treated with saline during adolescence, and saline during
adulthood. Adolescent THC exposure prevented this morphine-induced anxiety. One-way ANOVA p =
0.0006; pairwise t-tests, vehicle/saline vs. vehicle/morphine 0.93 vs. 3.67, p = 0.0048; vehicle/saline vs.
THC/morphine 0.93 vs. 0.50, p = 0.83; vehicle/morphine vs. THC/morphine 3.67 vs. 0.50, p = 0.006.
H, Schematic of the von Frey task of mechanical hypersensitivity.
I, Mice given saline vs. THC during adolescence showed no difference in baseline mechanical sensitivity.
Vehicle/morphine vs. THC/morphine 0.85 g vs 0.86 g, p = 0.93. **J**, Repeated morphine administration
leads to a reduction in the 50% mechanical threshold, indicative of pain hypersensitivity relative to mice
treated with saline during adolescence, and saline during adulthood. Adolescent THC exposure blunted
this morphine-induced pain hypersensitivity. One-way ANOVA p = 0.0032; pairwise t-tests, vehicle/saline
vs. vehicle/morphine 1.09 vs. 0.42, p = 0.0027; vehicle/saline vs. THC/morphine 1.09 vs. 0.69, p = 0.052;
vehicle/morphine vs. THC/morphine 0.42 vs. 0.69, p = 0.30.

Figure 3: **Mice treated with THC during adolescence show an elevation in drug-induced**

reinstatement of morphine-seeking behaviors later in life.

A, Timeline of experiments.

B, Time spent in the morphine-paired chamber during CPP pre-test, post-test, extinction, and

reinstatement. Mice treated with THC or vehicle during adolescence showed a similar trajectory except

for during reinstatement using a 5 mg/kg dose of morphine, at which time THC-treated mice showed an

elevation in reinstatement behavior.

C, Quantification of the time spent in the morphine-paired chamber in the reinstatement task relative to

the final extinction day. Vehicle/morphine-treated mice 722 s vs 801.3 s, p = 0.43, n = 12 each;

THC/morphine treated mice 780.7 s vs. 1015 s, p = 0.027, n = 8 each.

Figure 4: Brain-wide activity patterns in adolescent vehicle- and THC-treated mice following a reinstatement dose of morphine.

A, Representative sagittal section of a cFos-stained brain of an animal treated with THC during adolescence and following a reinstatement dose of morphine. Scale, 2 mm.

B, More cFos-expressing cells were detected in adolescent THC-treated mice than vehicle-treated mice following a dose of 5 mg/kg morphine (548,481 vs. 325,789 cells, p = 0.0051).

C, The top ten brain regions showing the highest-fold elevation in cFos labeling in adolescent THC-treated relative to vehicle-treated mice and the four regions showing an elevation in vehicle-treated mice. Groups are separated by the dashed red line. Shown are brain regions with an average > 150 cFos+ cells in each brain region.

D, Representative sagittal images of the GPe in adolescent vehicle-treated vs. THC-treated mice. Scale, 500 μm.

E, Anatomical connectivity matrix of the mouse brain, broken into four communities based on the Louvain method for community detection

F, Log fold-change in cFos labeling in each network community relative to one another (One-way ANOVA
p = 0.0002; purple (1) vs. blue (2) 0.68 vs. 0.75, p = 0.76; purple (1) vs. red (3) 0.68 vs. 0.61, p = 0.82;
purple (1) vs. yellow (4) 0.68 vs. 0.38, p = 0.0024; blue (2) vs. red (3) 0.75 vs. 0.61, p = 0.31; blue (2)
vs. yellow (4) 0.68 vs. 0.38, p = 0.0002; red (3) vs. yellow (4) 0.61 vs. 0.38, p = 0.05.
G, UMAP embedding of regions based on their correlated activity in adolescent vehicle-treated mice.
Each point is a brain region.
H, UMAP embedding of regions based on their correlated brain activity in adolescent THC-treated mice.
The regions are overall more tightly clustered than in panel G, suggesting a greater level of correlated
activity in adolescent THC-treated mice.
I, Hierarchical clustering of regions based on the cFos correlation matrix in adolescent vehicle-treated
mice. The VTA is within a separate activity-defined cluster from the frontal cortex regions of interest
(prelimbic cortex, PL; infralimbic cortex, ILA; anterior insula cortex, AI; anterior insula cortex, ventral part,
AIv; ACAd, anterior cingulate cortex, dorsal part; ACAv, anterior cingulate cortex, ventral part).
J, Hierarchical clustering of regions based on the cFos correlation matrix in adolescent THC-treated mice.
The VTA co-clusters with regions of the anterior cortex in these mice.
K, The number of hierarchical clusters in adolescent vehicle-treated and THC-treated mice across a
range of heights for cluster cut-offs in hierarchical trees. The THC-treated mice show reduced modularity
across a range of cutoff values.

Figure 5: THC-treated mice show a different brain-wide pattern relative to animals treated with

psychostimulants, opioids, or nicotine.

A, Schematic of experiments.

B, Representative image of the ventral midbrain of a DAT-Cre mouse showing starter cells in the VTA
and medial SNc. Scale, 1 mm.

C, Representative images of input cell populations in several brain sites, including the anterior cortex,
NAcMed, NAcCore, NAcLat, MHb, LHb, and PBN. Scale, 250 μm.

D, Bar graph plot showing the percentage of RABV-labeled inputs in control vs. THC-treated mice. Two-
way ANOVA, interaction p = 0.0062. Only the anterior cortex was significantly different following
correction for multiple comparisons (corrected p-value p = 0.029.

E, Schematic and explanation for how PCA analysis was carried out on the RABV tracing dataset.

F, Percent of cumulative variance explained by each principal component.

G, Heatmap of the contributions of each brain region, or feature, in the data to PCs 1 through 3.

H, Plot of PC1 and PC2 from brains of control and THC-treated mice.

I, Plot of PC1 and PC3 from brains of control and THC-treated mice.

J-L, Comparison of total PC values for each brain along **J** PC 1, **K** PC 2, and **L** PC 3. **J**, Control 3.3 vs.
THC -2.2, p = 0.0002. **K**, Control 0.13 vs. THC -0.09, p = 0.89. **L**, Control 0.57 vs. THC -0.38, p = 0.48.

M, To further quantify relationships between brains and conditions, the Euclidean distance between
each brain from control and THC-treated mice in the PC1 versus PC2 (or PC1 vs. PC3) coordinate
space was calculated. These distances were then plotted in a heatmap with brains being organized
by similarity assessed by hierarchical clustering.

N, Heatmap of the contributions of each brain region, or feature, in the data to PCs 1 through 3 in the
data presented in panels N-Q.

O, Plot of PC1 and PC2 from brains of control and drug-treated mice.

P, Plot of PC1 and PC3 from brains of control and drug-treated mice.

Q-R, Comparison of total PC values for each brain along **Q** PC 2, and **R** PC 3. For PC 1, One-way ANOVA
p = 0.83. **Q**, One-way ANOVA p < 0.0001; pairwise t-tests and asterisks presented in the figure include
THC comparisons only, although statistics are based on comparisons between all groups. Saline vs. THC
-2.43 vs. 2.96, p < 0.0001; Fluoxetine vs. THC -1.55 vs. 2.96, p = 0.0004; K/X vs. THC -1.23 vs. 2.96 p
= 0.0024; Drugs vs. THC 0.19 vs. 2.96, p = 0.0051. **R**, One-way ANOVA p = 0.0004; Saline vs. THC 0.70
vs. 1.85, p = 0.63; Fluoxetine vs. THC 0.88 vs. 1.85, p = 0.71; K/X vs. THC 0.19 vs. 1.85 p = 0.27; Drugs
vs. THC -0.79 vs. 1.85, p = 0.0005.

S, The Euclidean distance between each psychoactive drug-treated and control brain in the PC1 versus
PC2 coordinate space was calculated, plotted in a heatmap, and brains were organized by similarity and
assessed by hierarchical clustering.

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