Neural circuit basis of adolescent THC-induced potentiation of opioid responses in adult mice

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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 impacted morphine-induced behaviors across several dimensions, including potentiating reward and paradoxically impairing the development of morphine reward. We then mapped the whole-brain response to a reinstatement dose of morphine, finding that adolescent THC administration led to increased 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 that has the potential to influence dopaminergic response to morphine administration during adulthood. Our study adds to the rich literature on the interaction between drugs of abuse 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 during adolescence 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 number as those who misuse psychoactive drugs such as cocaine and opioids[3]. With the trend of legalization across the globe enabling increased access to cannabis, it is critical to understand how adolescent cannabis use impacts the brain and behavior. 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 consumption of psychoactive 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 leads 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 influences the adult brain 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. Despite the goal to reduce the clinical usage of opioids, opioids remain the primary treatment for post-operative pain management. Unfortunately, approximately eighty percent of those who use heroin

began using opioids for pain management[21,22]. As opioid exposure is thus in some cases largely

unavoidable, it is important to understand how lifestyle factors such as adolescent THC usage influence

response to opioids. In fact, the intersection of THC and opioids has been a question of substantial

interest for many decades, in part due to the hypothesis that cannabis use may predispose individuals to

opioid use[23–25]. 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[26]. Cross-

sensitization of THC and morphine has been demonstrated[27], and adolescent THC exposure can alter

expression of genes related to the endogenous opioid system[10], providing further evidence that these

drugs may work via common mechanisms.

Here, we administer a moderate dose of THC (5 mg/kg) for 14 days starting at postnatal day 30 (PD30),

during adolescence. We then test a variety of opioid-induced behaviors by administering morphine later

in life and assess how adolescent THC administration influences these behaviors. We also assess how

adolescent THC administration influences brain-wide activity patterns following a reinstatement dose 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 ($22^{o}C$ +/- $2^{o}C$) and humidity (55-65%). Mice

received ad libitum access to food and water.

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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 composed of 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.
Open field boxes were outfitted with contexts to differentiate them from the open field test for anxiety.
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. Mice were excluded if data
from >1 day of the task were missing.

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 were
subject to exclusion if the time spent on one side during the CPP pretest was less than 600 seconds or
more than 1200 seconds.

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. Withdrawal tests were
not performed on one cohort of mice.

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 in 50% of animals[28].

cFos induction and immunohistochemistry: One hour following injection of 5 mg/kg morphine, 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[29]. 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. Accurate detection of cFos+ neurons was ensured by manual quality control checks for each

brain. ClearMap[30] was then used to align the imaged brains to the Allen Brain Reference Atlas, and to

quantify the cFos+ neurons in anatomically-defined brain regions.

Network Analysis: Anatomical projection data from the Allen Mouse Brain Connectivity Atlas was used to

construct a connectivity matrix among the brain regions in our analysis[31]. 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 dividing 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[32]. 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[33]. Regions were clustered using HDBSCAN,

which can identify hierarchical clusters in continuous data based on variations in density of the data[34].

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[35].

<u>Rabies virus circuit mapping</u>: Rabies virus (RABV) monosynaptic tracing studies were carried out as

previously described[19,36], 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[19,36].

<u>Viruses and injection coordinates</u>: 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[35].

<u>Quantification and Statistical Analysis</u>: Statistics for all 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, 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 reduces morphine-induced locomotion, with no impact on morphine
reward when given during young adulthood
We first exposed mice to a daily 5 mg/kg/day dose of THC during adolescence, here defined as postnatal
days 30-43 (Figure 1A). This is considered a moderate dose that is mildly aversive and anxiogenic[37].
Animals receiving daily injections of vehicle on equivalent days were used as controls. 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 may
impact morphine reward, we performed a two-dose morphine conditioned place preference (CPP) 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, whereas 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). We observed that adolescent THC exposure
blunted morphine-induced locomotion during the five-day morphine administration window (RM 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.
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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 an 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). 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 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 an elevated
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). 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.
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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.
Mice showed normal CPP (e.g., Figure 1D) and the majority extinguished normally following 2 extinction
sessions (Figure 3A-B). To reduce confounding factors while examining reinstatement, mice were
excluded that did not show normal CPP or CPP extinction. 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 mice in response to a sub-threshold dose of morphine in mice given vehicle during
adolescence.
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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 a more robust drug-primed reinstatement than vehicle-
treated mice, 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 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.
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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[31]. 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).
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Following this analysis, we wanted to assess if these pathway-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, clusters which
mapped relatively consistently onto the adolescent vehicle-treated mice data (Figure 4G-H), with some
increased spread in 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), suggesting 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[19,38], and input connectivity was assessed. RABV-labeled input cells were observed throughout the brain, as in previous studies (Figure 5B-C). Overall, we observed several quantitative differences, quantified 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 the anterior cortex, 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[38,39]. 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 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, principally

an increase in connectivity from the anterior cortex.

DISCUSSION:

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

consequences for brain-wide activity following reinstatement and midbrain dopamine cell connectivity.

We found that adolescent THC exposure influences morphine responses across a variety of behavioral

dimensions, including locomotion, withdrawal-induced anxiety-related behavior, pain hypersensitivity,

and reinstatement. This broad range of effects is consistent with findings from studies that have explored

the influence of THC on morphine-induced behavioral changes, and vice versa[40,41]. 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, which has the potential to alter the dopaminergic
response to morphine administration later in life. Our study thus adds to a rich literature on the subject
by implicating several discreet circuits as potential drivers of the long-lasting behavioral effects induced
by adolescent THC exposure on responses to opioids later in life.

While adolescent THC exposure enhances the behavioral response to a reinstatement dose of morphine,
consistent with adolescent THC potentiating behavioral responses to morphine, it is perhaps surprising
that THC exposure blunts morphine withdrawal-induced anxiety and pain. Interestingly, behavioral profile
is consistent with the behavioral responses observed following exposure of drugs of abuse during
adolescence[42–46], 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[47]. 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. 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 each 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[8,9,41]. 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 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[36]. 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[48,49]. Furthermore, we recently showed that the extent of input labeling is

dependent on the activity of input connections[19,20,50]. 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[38]. 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 a drug of abuse. 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 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[19,20,50], with this signature being relatively similar regardless of the drug administered[19,38],

suggesting that these drugs induce similar changes in connectivity. Interestingly, in those studies we did

not observe an increase in inputs from the anterior cortex[19,20,38,50]. In fact, inputs from the anterior cortex

tend to be depressed following drug administration[19], and this reduction was one of the three brain regions

that showed a significant change in connectivity following drug administration. The anterior cortex,

particularly the mPFC, is thought to play important roles in the later stages of substance use disorder[51–

54]. The reduction in connectivity following a single dose, and increase in connectivity following repeated

doses, is thus consistent with that hypothesis of increased mPFC engagement later in 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 $p = 0.75$.

Vecchio/saline-treated mice are shown for comparison but were not included in statistical tests. Two-way

ANOA 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 vehicle 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 reinstatement 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 regions.

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.

H, UMAP embedding of regions based on their correlated brain activity in adolescent THC-treated mice.

I, Hierarchical clustering of regions based on the cFos correlation matrix in adolescent vehicle-treated

mice.

J, Hierarchical clustering of regions based on the cFos correlation matrix in adolescent THC-treated

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.

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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