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## **Behavioral Deficits and Associated Alterations in the Proteome in the Amygdala of Adolescent Rats Exposed to Delta9-Tetrahydrocannabinol as Juveniles**

Afzaal Nadeem Mohammed

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Behavioral deficits and associated alterations in the Proteome in the Amygdala of  
adolescent rats exposed to delta9-tetrahydrocannabinol as juveniles

By

Afzaal Nadeem Mohammed

A Dissertation  
Submitted to the Faculty of  
Mississippi State University  
in Partial Fulfillment of the Requirements  
for the Degree of Doctor of Philosophy  
in Environmental Toxicology  
in the College of Veterinary Medicine

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2018

Behavioral deficits and associated alterations in the Proteome in the Amygdala of  
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By

Afzaal Nadeem Mohammed

Approved:

---

Russell Carr  
(Major Professor)

---

Bindu Nanduri  
(Co-Major Professor)

---

Barbara L. Kaplan  
(Committee Member)

---

Jeffery B. Eells  
(Committee Member)

---

Mathew K. Ross  
(Committee Member)

---

Mark L. Lawrence  
Dean  
College of Veterinary Medicine

Name: Afzaal Nadeem Mohammed

Date of Degree: December 14, 2018

Institution: Mississippi State University

Major Field: Environmental Toxicology

Major Professor: Russell Carr

Title of Study: Behavioral deficits and associated alterations in the Proteome in the Amygdala of adolescent rats exposed to delta9-tetrahydrocannabinol as juveniles

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Candidate for Degree of Doctor of Philosophy

Abstract:  $\Delta$ 9-tetrahydrocannabinol ( $\Delta$ 9-THC) is an active component of marijuana. During recent years, the popularity of marijuana in the United States has increased tremendously. Marijuana edibles are a form of marijuana that has become very popular in recent years. These are easily accessible not only to adolescents but also to young children. According to recent statistical data, the consumption of marijuana edibles by children below the age of 5 has increased 600% in the states that have legalized marijuana. This has led to an adverse impact on children's health as evidenced by a sudden increase in the number of children seeking emergency assistance in hospitals. In the current research, we addressed the issue of possible persistent effects on children's behavior due to an earlier exposure to  $\Delta$ 9-THC. Juvenile rats were treated with 10 mg/kg of  $\Delta$ 9-THC from postnatal day 10 through 16. Once they reached adolescence, these rats were tested using several behavioral paradigms. To evaluate the biological basis for the behavioral deficits observed, brain samples obtained from these rats were subjected to proteomic analysis to determine any altered pathways related to the behavior. Our behavioral data indicated that juvenile exposure to  $\Delta$ 9-THC has no effect on anxiety-

related behavior in adolescents. However, we observed a significant effect of treatment on multiple parameters related to social interactions. Of these, episodes and time of social play were significantly increased in the  $\Delta 9$ -THC treated rats suggesting alterations in the reward circuit function occurring as a result of developmental  $\Delta 9$ -THC exposure. In the proteomics, we observed a significant effect on relevant canonical pathways such as the changes in thrombin and opioid signaling. Thrombin signaling in neurons is associated with processes involved in the connection between neurons and opioid signaling is involved in the activation processes of the reward circuit suggesting that juvenile  $\Delta 9$ -THC exposure alters these processes in adolescence which could have detrimental effects on behavior. Overall, our data suggest that consumption of edibles by juveniles leads to altered behavioral and biochemical outcomes in adolescence. This may be detrimental in terms of the appropriate acquisition of skills necessary for meeting the challenges in future life.

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## TABLE OF CONTENTS

ACKNOWLEDGEMENTS.....	ii
LIST OF TABLES.....	vii
LIST OF FIGURES.....	ix
CHAPTER.....	1
I. DELTA9-TETRAHYDROCANNABINOL: MOLECULAR TARGETS, EFFECTS ON EMOTIONAL AND SOCIAL BEHAVIOR AND INCREASING EXPOSURE THROUGH MARIJUANA EDIBLES.....	1
1.1 Background.....	1
1.2 The Endocannabinoid System.....	8
1.2.1 Functions of CB1 receptors in developing brain.....	9
1.2.2 Mechanism of CB1 receptor function.....	12
1.2.3 Function of CB1 receptor in mediating behavior.....	14
1.3 Review of Literature.....	18
1.3.1 Elevated plus maze (EPM).....	18
1.3.2 Social play.....	23
1.3.2.1 Social play in developmental disorders.....	24
1.3.2.2 Structure and neurobiology of play.....	24
1.3.2.3 Effect of $\Delta$ 9-THC on play behavior.....	26
1.4 Proteomic Approach to Study Behavior.....	27
1.5 Reference:.....	30
II. EFFECT OF REPEATED JUVENILE EXPOSURE TO $\Delta$ 9- TETRAHYDROCANNABINOL ON ANXIETY-RELATED BEHAVIORS AND SOCIAL INTERACTIONS IN ADOLESCENT RATS.....	43
2.1 Introduction.....	43
2.2 Materials and Methods.....	46
2.2.1 Chemicals.....	46
2.2.2 Animals.....	47
2.2.3 Exposure.....	47
2.2.4 Elevated plus maze (EPM).....	47
2.2.5 Social behavior (PND37-38).....	49

2.2.6	Statistical analysis.....	50
2.3	Results.....	51
2.3.1	Weight gain .....	51
2.3.2	Elevated plus maze .....	51
2.3.3	Open arm activity.....	52
2.3.4	Ethological oriented parameters and overall activity.....	53
2.3.5	Social behavior .....	55
2.4	Discussion .....	60
2.5	Reference:.....	67
III.	ALTERATIONS IN THE PROTEOMIC PROFILE AND CANONICAL PATHWAYS IN ADOLESCENT RATS EXPOSED TO $\Delta$ 9-TETRAHYDROCANNABINOL AS JUVENILES.....	76
3.1	Introduction .....	76
3.2	Materials and Method .....	78
3.2.1	Chemicals .....	78
3.2.2	Animals .....	78
3.2.3	Exposure.....	79
3.2.4	Proteomic analysis .....	79
3.2.5	Data analysis.....	81
3.2.6	IPA and gene ontology.....	81
3.2.7	Quantitative PCR (QPCR).....	82
3.3	Results and Discussion .....	84
3.3.2	Gaq signaling.....	90
3.3.3	Opioid signaling.....	91
3.3.4	Rho GTPase and thrombin signaling .....	92
3.4	Conclusion.....	93
3.5	Reference:.....	99
IV.	ALTERATIONS IN CB1-RELATED CANONICAL PATHWAYS DURING SOCIAL PLAY IN ADOLESCENT RATS REPEATEDLY EXPOSED TO $\Delta$ 9-TETRAHYDROCANNABINOL AS JUVENILES.....	106
4.1	Introduction .....	106
4.2	Materials and Method .....	108
4.2.1	Chemical.....	108
4.2.2	Animals .....	108
4.2.3	Exposure.....	109
4.2.4	Animal handling .....	109
4.2.4.1	Behavior rats.....	109
4.2.4.2	Non-behavior rats.....	110
4.2.5	Proteomic analysis .....	110
4.2.6	Data analysis.....	112
4.2.7	IPA and gene ontology.....	113
4.2.8	Quantitative PCR (QPCR).....	113

4.3	Results and Discussion .....	115
4.3.1	Control behavioral verses $\Delta$ 9-THC-treated behavioral comparison .....	116
4.3.2	$\Delta$ 9-THC-treated behavioral versus $\Delta$ 9-THC-treated non- behavioral comparison .....	123
4.3.3	Control behavior and control non-behavior comparison.....	131
4.4	Conclusion.....	138
4.5	Reference:.....	139
V.	CONCLUSION.....	146
5.1	Reference:.....	151
A.	Lists of Proteins Obtained from Proteomic Analysis Under Different Experimental Conditions .....	153

## LIST OF TABLES

1.1	Specific behaviors represented by level of exploration performed by rats or mice in different arms of elevated plus maze.....	19
1.2	Different types of activities performed during the act of play and associated behaviors represented by each activity .....	26
3.1	Primers, probes, and assay names for reference and target genes .....	83
3.2	Canonical pathways predicted using differentially expressed proteins in the amygdala of adolescent rats (PND38) exposed to either corn oil or 10 mg/kg $\Delta$ 9-THC during juvenile period (PND10-16).....	86
3.3	List of differentially expressed protein involved in the relevant canonical pathways shown in figure 3.3.....	97
4.1	Primers, probes, and assay names for reference and target genes .....	114
4.2	List of canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats receiving either corn oil or 10 mg/kg $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play.....	119
4.3	Differentially expressed proteins involved in the relevant canonical pathways shown in Figure 4.2.....	121
4.4	List of canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats receiving only 10 mg/kg $\Delta$ 9-THC during juvenile ages. Tissue was extracted on PND38, either 3-hours following play (behavior $\Delta$ 9-THC) or without being subjected to play (non-behavior $\Delta$ 9-THC). .....	126
4.5	Differentially expressed proteins involved in relevant canonical pathways shown in figure 4.5 .....	129
4.6	List of canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats exposed to corn oil during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior control) or without being subjected to play (non-behavior control). .....	133

4.7	List of differentially expressed proteins involved in canonical pathways listed in figure 4.....	136
A.1	List of differentially expressed proteins obtained from vehicle or $\Delta 9$ -THC treated non-behavior rats.....	154
A.2	List of differentially expressed proteins obtained from vehicle or $\Delta 9$ -THC treated behavior rats.....	157
A.3	List of differentially expressed proteins obtained from $\Delta 9$ -THC treated behavior or non-behavior rats.....	158
A.4	List of differentially expressed proteins obtained from $\Delta 9$ -THC treated behavior or non-behavior rats.....	160

## LIST OF FIGURES

- 2.1 Plot of weight gain of male and female rat pups exposed daily to either corn oil (control) or 10 mg/kg of  $\Delta$ 9-THC as juveniles from postnatal days 10-16. Values are expressed as weight  $\pm$  SE (n = 31-37). For both sexes, weight gain in treated rats was significantly different from that in the controls ( $p \leq 0.05$ ) at all time points..... 56
- 2.2 Open arm activity in the elevated plus maze under different levels of illumination as described in the Materials and Methods. Number of entries into open arms under high (A) and low (B) illumination, percent entries into open arms under high (C) and low (D) illumination, time spent in the open arms under high (E) and low (F) illumination, and percent time spent in the open arms under high (G) and low (H) illumination, in male and female adolescent rats (PND29) exposed to either corn oil or 10 mg/kg of  $\Delta$ 9-THC as juveniles from postnatal days 10-16. Values are expressed as means  $\pm$  SEM or percent  $\pm$  SEM (n=11-30). Brackets with asterisks (\*) are statistically significant ( $p \leq 0.05$ ) from one another ..... 57
- 2.3 Open arm activity in the elevated plus maze under different levels of illumination as described in the Materials and Methods. Number of entries into open arms under high (A) and low (B) illumination, percent entries into open arms under high (C) and low (D) illumination, time spent in the open arms under high (E) and low (F) illumination, and percent time spent in the open arms under high (G) and low (H) illumination, in male and female adolescent rats (PND29) exposed to either corn oil or 10 mg/kg of  $\Delta$ 9-THC as juveniles from postnatal days 10-16. Values are expressed as means  $\pm$  SEM or percent  $\pm$  SEM (n=11-30). Brackets with asterisks (\*) are statistically significant ( $p \leq 0.05$ ) from one another ..... 58
- 2.4 Social behavior interactions as described in the Materials and Methods. Frequency of social exploration (A), frequency of nape attacks (B), frequency of play fighting (C), and time spent play fighting (D) in adolescent rats (PND38) exposed to either corn oil or 10 mg/kg of  $\Delta$ 9-THC as juveniles from postnatal days 10-16. Insets are values separated by sex. Values are expressed as means  $\pm$  SEM (n=10-12). Bars with asterisks (\*) are statistically significant ( $p \leq 0.05$ ) from controls. .... 59

3.1	Network overlap generated by IPA by list of differentially expressed proteins at fishers exact P-value of 0.05 (A) and 0.08 (B). .....	94
3.2	GO term analysis of differentially expressed proteins obtained from amygdala of adolescent rats (PND38) repeatedly exposed to either corn oil or 10 mg/kg $\Delta$ 9-THC during juvenile ages (PND10-16).....	95
3.3	Canonical pathways associated with differentially expressed proteins obtained from amygdala (PND38) of rats exposed repeatedly to either corn oil or 10 mg/kg $\Delta$ 9-THC at juvenile ages (PND10-16). .....	96
3.4	Change in mRNA expression in amygdala of adolescent rats (PND38) repeatedly exposed to either corn oil or 10 mg/kg of $\Delta$ 9-THC during juvenile ages (PND10-16).....	98
4.1	GO term analysis of differentially expressed proteins obtained from amygdala tissue of behavioral rats repeatedly exposed to either corn oil or 10 mg/kg $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted 3-hours following social play on PND38. ....	120
4.2	Relevant canonical pathways obtained from IPA analysis of differentially expressed proteins obtained from amygdala tissue of behavioral rats repeatedly exposed to either corn oil of 10 mg/kg $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted 3-hours following social play on PND 38. ....	121
4.3	Relative change in mRNA expression of G protein subunits in amygdala of behavioral rats repeatedly exposed to either corn oil or 10 mg/kg of $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted 3-hours following social play on PND38. ....	123
4.4	GO term analysis of differentially expressed proteins obtained from amygdala of adolescent rats repeatedly exposed to only $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior $\Delta$ 9-THC) or without being subjected to play (non-behavior $\Delta$ 9-THC).....	125
4.5	Relevant canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats receiving only 10 mg/kg $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior $\Delta$ 9-THC) or without being subjected to play (non-behavior $\Delta$ 9-THC). ....	128

4.6	GO term analysis of differentially expressed proteins obtained from amygdala tissue of rats repeatedly exposed to corn oil during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior controls) or without being subjected to play (non-behavior controls). ....	133
4.7	Relevant canonical pathways obtained from differentially expressed proteins obtained from amygdala of adolescent rats repeatedly exposed to only corn oil during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior controls) or without being subjected to play (non-behavior controls). ....	135



CHAPTER I  
DELTA9-TETRAHYDROCANNABINOL: MOLECULAR TARGETS, EFFECTS ON  
EMOTIONAL AND SOCIAL BEHAVIOR AND INCREASING EXPOSURE  
THROUGH MARIJUANA EDIBLES.

**1.1 Background**

Marijuana refers to the dried leaves, flowers, seeds or stems of plants known as *Cannabis sativa* or *Cannabis indica*. Marijuana resins can be obtained from these plants in the form of hash oil, wax, budder, or shatter which are widely smoked by the adolescent and adult population in order to get ‘high’ (NIDA: Marijuana facts 2017). Even though marijuana is currently among the most widely used illicit drug, it has been historically used for many commercial and medicinal purposes. Marijuana plants were commonly grown in the 17<sup>th</sup> century for the production of rope, sails and clothing. The medical applications for marijuana originated from Chinese folk medicine that included the use of marijuana for a variety of treatments as early as the 27<sup>th</sup> century BC (McKenna 2014). In the United States, the use of marijuana for medical and recreational purposes increased in the early 19<sup>th</sup> century and marijuana was an active ingredient in more than 20 drugs. During the same period, the prohibitionist attitudes towards drugs was increasing and marijuana was perceived as a dangerous drug able to induce sudden violent and murderous actions in intoxicated individuals (Carliner et al., 2017). In 1937, the federal Marijuana Tax Act was

passed which prohibited the use of marijuana for recreational purposes. Since this law did not ban medical uses, marijuana continued to be used for medicinal purposes under this act. However, this law discouraged the use of marijuana for medical purposes by placing the burden of detailed reporting on physicians who were in favor of prescribing marijuana (McKenna 2014). According to the Control Substance Act of 1970, the United States Drug Enforcement Administration (DEA) categorized marijuana as schedule 1 drug (i.e. a drug with no medical application and high potential of abuse). This categorization made both recreational and medical use of marijuana illegal. Despite the federal stance against marijuana use, many Americans are in favor of marijuana being used for medical purposes. Owing to the favorable attitude of public towards marijuana, states across the nation have been eliminating the legal restrictions against marijuana. California was the first state to allow marijuana for medical uses in 1996. Following that, more states allowed marijuana for medical purposes. These changes have had consistent public support due to the inclination of the majority of the population to live in jurisdictions where marijuana is legal. By 2017, 28 states have legalized marijuana for medical purposes and 7 of them allow recreational use as well (Carliner et al., 2017; Murphy and Carnevale 2016).

While legalization has alleviated the restrictions against using marijuana for medical purposes, it has also increased voluntarily self-intoxication among adolescents (Hopfer 2014). This increase in self-intoxication can be attributed not only to the increased ease of availability but also to the positive or permissible attitudes and diminished perception of marijuana-associated risks of adolescents (Walker et al., 2017). Adolescents appear to be unaware of the neurological and/or psychotic problems associated with the early-onset use of marijuana as reported by number of studies. According to the Diagnostic and Statistical

Manual of Mental Disorders, 4<sup>th</sup> edition (DSM-IV), abuse and dependence are two major problems associated with marijuana and account for 75-80% of total marijuana use disorder cases. In a statistical study, marijuana abuse and dependence cases among the US adult population occurring during 1991-1992 were compared with those occurring during 2001-2002. There was a significant increase in the level of abuse and dependence cases in the latter period despite no significant increase in marijuana use among the adult population. However, the comparatively higher potency of marijuana available in the latter period might be an additional factor in the increased level of disorders (Compton et al., 2004). Another consequence of marijuana legalization has been an increase in the popularity of marijuana among school children. A study in Monitor the Future (MTF) reported that marijuana smoking has been a common practice among 12<sup>th</sup> graders but since the mid-1990s, it has become increasingly popular among 10<sup>th</sup> graders and 8<sup>th</sup> graders because of the lowered perceived risk and increase in the ease of availability during that period (MTF 2015).

Early onset of marijuana use is associated with certain structural abnormalities in the brain that eventually lead to behavioral deficits. Neuronal pruning is one of the essential processes by which the developing brain forms into a mature brain by establishing proper connections across the brain regions. Disrupted pruning can lead to mental disorders such as autism, schizophrenia, depression and others (Boksa 2012). Neuroimaging studies have reported that neuronal pruning can be drastically affected in cannabis users (Filbey et al., 2015). In addition to neuronal pruning, heavy cannabis consumption is also related to a reduction in the area of white matter that is associated with increased impulsivity (Gruber et al., 2014), and increased activity in cortical areas that is associated with attention

difficulties (Abdullaev et al., 2010). Currently, the available literature is limited to addressing the adverse effects of marijuana use in adolescents. Despite the documented increase in the availability of and exposure to marijuana in children, there have been no studies that have addressed the long-term effects of marijuana exposure in children.

Edible marijuana is one of the choice methods for marijuana intake along with smoking and vaping. Starting in 2014, Colorado allowed the commercial production of edible marijuana and marijuana is now available in a form that mimics popular candies or sweets. These products can be legally purchased in retail stores in four states; Colorado, Washington State, Alaska, and Oregon (MacCoun et al., 2015). In California, marijuana edibles are available in dispensaries but only those who possess a medical marijuana card which can be obtained by anyone including those below 18 years as long as they are under a parent's supervision (Borodovsky et al., 2016). According to sales (NIDA 2014) and social networking surveys (Borodovsky et al., 2016; Lamy et al., 2016), the popularity of marijuana edible products is increasing in states with legalized marijuana as compared to non-legalized states. This has created serious health concerns because of the enhanced risk of edible marijuana being consumed by children even though current laws restrict recreational marijuana use to those under the age of 21 (MacCoun et al., 2015). In addition, the packaging of these edibles in bright colors is more likely to attract children who are unaware of the risk. This 'child friendly' packaging is increasing either deliberate or unintentional consumption of marijuana among children and there has been increased reports of cases of childhood intoxication especially in states with legalized marijuana (MacCoun et al., 2015). The data from poison control centers indicates higher rates of emergency calls and an increase in the number of patients seeking care at children's

hospitals in states with legalized marijuana because of the unintentional ingestion of marijuana in children of age below 9. Soon after the commercialization of marijuana edibles in Colorado in 2014, there was 70% increase in the calls related to marijuana exposure compared to the prior year (Ghosh et al., 2013). In another study conducted at a tertiary-care children's hospital in Colorado, there were 14 cases, all below the age of 12, that involved intoxication related to marijuana ingestion during 2009-2011 as compared to none during 2005-2009 (Wang et al., 2013). Marijuana edibles were originally introduced for medical purpose and meant for medical-prescribing to those who are not comfortable with smoking marijuana or children who can take them under caregiver's supervision. However, this has led to unintentional unwanted childhood exposure.

The main psychoactive component of marijuana is  $\Delta^9$ -tetrahydrocannabinol commonly known as  $\Delta^9$ -THC. When marijuana is consumed through any route of exposure, the concentration of  $\Delta^9$ -THC increases in the blood and is mainly responsible for the marijuana-related effects (NIDA 2017).  $\Delta^9$ -THC was first isolated in 1964 (Gaoni and Mechoulam 1964) and has been frequently studied for its pharmacological and psychoactive effects. In rodents,  $\Delta^9$ -THC is known to produce myriad of behavioral effects of which the most common ones are catalepsy, analgesia, hypothermia and hypolocomotion. These effects are commonly known as 'tetrad model' of cannabimimetic activity (Martin et al., 1991). Catalepsy is muscular rigidity or fixity of posture regardless of external stimuli. Several studies reported that  $\Delta^9$ -THC could induce catalepsy in rodent models by using bar catalepsy test (Marchese et al., 2003; Monory et al., 2007; Sano et al., 2008). Analgesia is reduced response to painful stimuli and has been detected following  $\Delta^9$ -THC exposure by measuring reaction time using the hot plate analgesia test (Monory

et al., 2007) or the tail-flick latency test (Fuentes et al., 1999). Similarly,  $\Delta 9$ -THC exposure is also associated with reductions in body temperature and locomotor activity as detected by measuring the rectal temperature and performance in the open field test, respectively (Monory et al., 2007).

To relate the route of exposure and the change in  $\Delta 9$ -THC levels in the blood, Lemberger and colleagues (1972) compared the plasma concentration of  $\Delta 9$ -THC following intravenous, ingestion, or inhalation administration of marijuana. The plasma  $\Delta 9$ -THC level was at its peak 70 min following inhalation and 3 hours following ingestion. Ingestion could be more dangerous route of exposure because of increased  $\Delta 9$ -THC bioavailability since it does not require combustion, which is the case for inhalation. Around 50% of  $\Delta 9$ -THC is lost in the burning process and the amount of uptake of the remainder depends upon the lung capacity and the amount of time that the smoker retains the smoke inside the lungs before exhalation (Elzinga et al., 2015). The onset of effects is rapid following inhalation. However, the onset of effects following ingestion is much slower than smoking because of delay of entrance of  $\Delta 9$ -THC into the blood stream. This factor is a prominent cause of marijuana-related overdoses because of the misinterpretation that initial dose did not induce the desired effect and the continued intake of more of the edible marijuana product (Brodovsky et al., 2016).

The  $\Delta 9$ -THC content in marijuana edibles is another major concern that could be related to the increase in intoxications. Over the time course of marijuana legalization, the  $\Delta 9$ -THC content in the products has increased to more than 20% from 15% (Ghosh et al., 2015). Often, underestimation of the  $\Delta 9$ -THC content on labeling is worsening this

problem. In a sample of 75 marijuana edibles collected from four different states, more than half of these edibles contained a label that underestimated the  $\Delta$ 9-THC content of the product (Vandrey et al., 2015). This could be a significant contributor towards overdose. The  $\Delta$ 9-THC content of marijuana edibles are calculated by number of  $\Delta$ 9-THC servings present in the edibles. Considering the increasing intoxication cases related to  $\Delta$ 9-THC ingestion, states that allow edible sales are implementing the restrictions on the amount of  $\Delta$ 9-THC in each serving.  $\Delta$ 9-THC content is set to be 10 mg/serving in Colorado and Washington State (MacCoun et al., 2015), and 5 mg/serving in Alaska and Oregon (Gourdet et al., 2017). However, there is not a policy that instructs manufacturer to limit the number of servings in each edible or to include on the label, the maximum daily intake of number of servings and/or edibles (Ghosh et al., 2015).

The primary target of  $\Delta$ 9-THC in the brain is the type I cannabinoid receptors (CB1; Devane 1988; Belue et al., 1994) which is a component of the endocannabinoid system. Most of the behavioral effects induced by  $\Delta$ 9-THC in laboratory studies are due to its action on CB1 receptors. The behavioral effects induced by  $\Delta$ 9-THC are because of its detrimental effects on developing neurons. Tyrosine Hydroxylase (TH) is an enzyme responsible for maturation of developing dopaminergic neurons (Fernandez-Ruiz et al., 1999). Gestational exposure to  $\Delta$ 9-THC has been reported to reduce the mRNA level of TH during the gestational period (Bonin et al., 1995), the postnatal/juvenile period, and the adult stage of life in rats (Suarez et al., 2000). Cholecystokinin is involved in the growth cone formation and axonal migration in developing neurons and prenatal  $\Delta$ 9-THC exposure reduces the density of cholecystokinin containing interneurons in hippocampus of juvenile rats (Vargish et al., 2017). Synapsin is a synaptic protein present in the axonal terminal and

important in establishing synaptic functions and exposure to  $\Delta 9$ -THC is associated with reduction in the mRNA levels of synapsin and CB1 receptor in adolescent rats (Steel et al., 2011). Exposure to  $\Delta 9$ -THC during adolescence is also related to hypofunction of GABA neurotransmission in prefrontal cortex of adult rats and this decrease in GABA neurotransmission was accompanied by an increase in the dopaminergic neurotransmission in ventral tegmental area (Renard et al., 2017).

## **1.2 The Endocannabinoid System**

The endocannabinoids are a group of neuro-modulatory lipids involved in a variety of physiological processes such as appetite, mood, sensation, and memory. These are not stored in vesicles but are produced through an on-demand basis when neurons are in excited state. This excitation of neurons leads to increase in intracellular calcium. The increase in intracellular calcium activates calcium-dependent N-acetyl transferase (NAT) which catalyzes the rate limiting step in the production of anandamide (Kano et al., 2009). This leads to increased production of the endocannabinoids that are then released from neurons and bind to the CB1 receptors (Lutz 2004). The mechanism by which the endocannabinoids exert their function was described by Alger and Marty (2001) who studied cerebellar purkinje neurons and hippocampal neurons. They observed that depolarization in the postsynaptic GABA cell leads to release of retrograde messengers that travel to presynaptic locus to suppress the release of neurotransmitter in that neuron. They called it “depolarization induced suppression of inhibition” or DSI. They also observed that this retrograde messenger was performing an analogous function in glutamatergic neurons by inhibition of excitation which they called as “depolarization induced suppression of excitation” or DSE. Both DSI and DSE can be blocked by CB1



receptor antagonists and enhanced by CB1 agonists indicating that the retrograde messenger inducing DSE and DSI is an endogenous cannabinoid or an endocannabinoid (Alger 2004). There are two main endogenous lipids, which bind to the CB1 receptors that are Arachidonoyl ethanolamine or Anandamide (AEA), and 2-Arachidonoyl glycerol (2-AG). These are produced in the postsynaptic neuron, cross the synapse, and stimulate CB1 receptors that are present on the presynaptic neuronal membrane. After stimulating the CB1 receptor, these lipids are taken up into the neurons through specific reuptake transporters and then degraded. The enzymes fatty acid amid hydrolase (FAAH) degrades anandamide and monoacylglycerol lipase (MAGL) degrades 2-AG (Kano et al., 2009).

### **1.2.1 Functions of CB1 receptors in developing brain**

Being a part of the endocannabinoid system, the CB1 receptor plays a vital role in brain development. Receptor dimerization is a critical phenomenon during neuronal development. Diverse receptors, such as the neurotrophic and growth factor receptors, are recruited to the CB1 receptor which implicates a role for the CB1 receptor in neuronal development (Harkany et al., 2008). Gestational day (GD) 11-14 is an important period in rats because this is the period where formation of the neural tube from ectodermal cells occurs. The brain and spinal cord are subsequently differentiated from the neural tube (Semple et al., 2013). CB1 mRNA is detected during this period suggesting that the cannabinoid receptors are important for these developmental processes. In addition to presence of CB1 mRNA, an increase in GTP-binding protein occurs in the presence of a cannabinoid agonist. This GTP-binding was mainly observed during the gestational period in brain regions where the CB1 receptors are scarcely distributed or practically absent in the adult brain. These brain regions are enriched in neuronal fibers and include white

matter areas of the midbrain and brainstem (Romero et al., 1997). Later, it was reported that in white matter areas, the CB1 receptors are located in the migrating axons whose cell bodies are located at significant distance from the site of the functioning receptor. This emphasizes the role of CB1 receptor in synaptogenesis (Fernandez-Ruiz et al., 2000). This role is also supported by the fact that the expression of the CB1 receptor is positively correlated with expression of Brain Derived Neurotropic Factor (BDNF), which is involved in axonal growth and migration during development (Maison et al., 2009).

Other studies have observed that activation of CB1 receptors by external cannabinoids during pregnancy can be detrimental to key proteins involved the brain development. Neural cell adhesion molecule (L1) is responsible for cell-cell and cell-matrix interactions allowing proteins to perform their functions such as cell proliferation, migration, synaptogenesis, and myelin formation. Mutation in the L1 genes leads to malformation of brain. Activation of the CB1 receptors by  $\Delta$ 9-THC exposure during pregnancy resulted in down regulation of genes responsible for L1 in rat brain during the fetal (Gomes et al., 2003) and early postnatal period (Gomes et al., 2007).

Irregular activation of CB1 receptors leading to abnormalities in the brain development is not only restricted to fetal period. Activation of the CB1 receptors as a result of exposure to  $\Delta$ 9-THC during the perinatal ages can also be harmful enough to cause certain behavioral and neurochemical abnormalities which remain into the later stages of life. For example, decreased locomotor activity and increased stereotypic behaviors such as self-grooming and shaking have been observed in  $\Delta$ 9-THC treated rats and these were accompanied by decreased presynaptic activity of niagro-striatal dopaminergic neurons, decreased striatal

dopamine D1 receptor intensity, and increased dopamine turnover (Novarro et al., 1994). Another long-term effect of  $\Delta 9$ -THC exposure is a change in the pattern of socio-sexual and social interactions. This change is accompanied by an increase in the turnover of mesolimbic dopamine (Novarro et al., 1996). Both of these studies involved adult rats who were exposed to hashish extracts during postnatal day (PND) 1-24. These studies establish a link between behavioral alterations in adults and the dopamine neurotransmitter system. Dopamine is mainly produced by the action of the enzyme tyrosine hydroxylase (TH). The CB1 receptors are present in the neurons containing TH and activation of the CB1 receptors is associated with an increase in the level and activity of TH. Interestingly, the CB1 receptors were detected in neurons containing TH during developmental stage and disappear from these neurons in adult brain (Hernandez et al., 2000). TH contributes to many different aspects of brain development such as differentiation of dopaminergic neurons, formation of connectivity, axonal guidance, and synaptogenesis (Fernandez-Ruiz et al., 1999). Therefore, behavioral abnormalities due to perinatal  $\Delta 9$ -THC exposure reported in the previously mentioned studies may be occurring because of effects of CB1 activation on TH.

The complexities of brain development are not limited to fetal and perinatal ages. The neuronal maturation and synaptogenesis continue to occur during the adolescent ages as evidenced by the reduction of grey matter. This reduction is mainly due to neuronal pruning, synaptogenesis (Whitford et al., 2007) and neuronal maturation of different neurotransmitter systems such as the endocannabinoid, GABA, dopaminergic, serotonergic, and adrenergic systems which mature to adult levels by pruning (Sturman and Moghaddam 2011). Exposure to  $\Delta 9$ -THC during adolescence is associated with certain

neurological disorders in later life such as long lasting cognitive deficits and impaired spatial and non-spatial learning (Cha et al., 2006). Interestingly, the learning deficits observed by Cha et al were specific to adolescent exposure whereas adults were more tolerant to the  $\Delta$ 9-THC induced learning impairments in a water maze task.

In light of the above studies concerning the CB1 receptors, it is obvious that there is a defined pattern of CB1 expression during brain development. Any alteration in this pattern due to exposure to exogenous substances like  $\Delta$ 9-THC may cause persistent changes in this brain development due to effects on certain neuronal activities which would consequently lead to adverse behavioral outcomes.

### **1.2.2 Mechanism of CB1 receptor function**

CB1 receptors are the most abundant G-protein coupled receptors in the brain and are mostly located on inhibitory neurons (GABA) and excitatory neurons (glutamate). They are differentially distributed among the different brain regions potentially because of the diverse functions that these receptors are involved in. Dense populations of the CB1 receptors were located in regions such as the basal ganglia, substantia nigra pars reticulata, globus pallidus (Herkenham et al. 1990). Later, using immunohistochemical analysis and ligands specific for CB1 receptors, the sub-cellular localization of CB1 receptors in the brain areas like hippocampus and amygdala was revealed that they are present on presynaptic GABA interneurons (Katona et al., 2001). As demonstrated by Hewlett and colleagues (1984),  $\Delta$ 9-THC-mediated CB1 activation leads to inhibition of adenylyl cyclase activity and a decrease in cAMP production. This inhibition was not observed in the presence of pertussis toxin which prevents the binding of  $G_{i/o}$  subunit to G-protein

couple receptors. This identified the role of Gi/o subunit in the functioning of the CB1 receptor. It has been demonstrated that both cannabinoid agonist-mediated inhibition of adenylyl cyclase and CB1-mediated adenylyl cyclase inhibition requires GTP. This suggests a pharmacological similarity in agonist-mediated and receptor-mediated action on adenylyl cyclase (Pacheo et al., 1991). This is further strengthened by the fact that agonist-mediated inhibition of adenylyl cyclase is blocked by selective CB1 antagonists (Rinaldi-Carmona et al., 1994).

Functioning of the CB1 receptors is not only mediated through Gi/o signaling and additional mechanism have been proposed involving binding of CB1 receptors to the Gs subunit instead of the Gi/o subunit (Hewlett et al., 2004, 2005). Binding of Gs to the G-protein receptor involves intracellular calcium signaling. One study reported that activation of the CB1 receptors following administration of an agonist inhibits the N-type and Q-type calcium channels. These channels normally allow the entry of calcium ions inside the neurons, increase in the calcium current, and facilitate the release of neurotransmitters stored in small vesicles in presynaptic neurons (Mackie and Hille 1992). It was also reported that agonist-mediated CB1 activation can cause stimulation of inwardly rectifying potassium channels and potassium current (Mackie et al., 1995). This study also confirmed that CB1-mediated effects on calcium and potassium channels were dependent on GTP-binding protein, but were not consequences of cAMP depletion, because the CB1-mediated inhibition of calcium channels was not reversed in presence of an external cAMP analogue (Mackie et al., 1995). Later, it was proposed that the cannabinoid-mediated effects on potassium and calcium channels occurs through binding of the G beta-gamma subunit which dissociated from the Gi subunit following CB1 activation (Ibsen et al., 2017). This

suggested the functioning of CB1 receptors through alternate subunits rather than only through Gi/o subunit. Several other studies support these findings in that CB1 receptors function by stimulating, rather than inhibiting the activity of adenylyl cyclase and increase the production of cAMP (Maneuf and Brotchie 1997; Glass and Felder 1997; Bonhaus et al., 1998; Felder et al., 1998). These studies observed that the CB1-mediated increase in adenylyl cyclase activity and cAMP accumulation could be due to the recruitment of the Gs subunit instead of the Gi/o subunit. These studies have observed that in the presence of pertussis toxin, CB1 activation can block the inhibition of adenylyl cyclase and increase cAMP accumulation that can be reversed by administration of CB1 antagonist (Felder et al., 1998). This stimulatory effect of the CB1 receptors on cAMP implies the participation of the Gs subunit. However, the high abundance of the Gi/o subunit lowers the affinity of the Gs subunit to bind to the CB1 receptor, thus making the stimulatory action less likely to occur (Glass and Felder, 1997). Another proposed mechanism of stimulatory action by the CB1 receptor is the association with specific isoforms of adenylyl cyclase in target tissues. While the activation of the CB1 can inhibit the 5/6 or 1/3/8 isoform families of adenylyl cyclase, it can also activate the 2/4/7 isoform family of adenylyl cyclase resulting in a stimulatory action (Rhee et al., 1998). Alternatively, the differential binding of subunits to CB1 receptors may be due to the different agonists used in previous studies. Different agonists can cause different conformational changes in CB1 receptors and these conformational changes dictate the binding of specific subunits (Ibsen et al., 2017).

### **1.2.3 Function of CB1 receptor in mediating behavior**

After the discovery that the CB1 receptor was a primary target of the cannabinoids, they have been studied for their role in memory, as well as in other types of behaviors. Stress is

one of the primary factors that plays a fundamental role in mental illnesses like depression and anxiety. When exposed to the same type of stress repeatedly, it leads to habituation which is a process of progressive decrease in the stress response after repeated exposure to same stress (Patel and Hillard, 2008). Stress habituation is controlled by glutamate release and has a prominent effect on memory encoding because it determines the ability to distinguish between stimuli that are worthy of memory consolidation and stimuli that are not. Stress habituation and memory are two interlinked processes that are controlled by the CB1 receptor through glutamate release (Litvin et al., 2013).

Social discrimination is a behavioral paradigm used to test social memory in animals. Social memory is predicated by the discriminative indices such as amount of time the animal spends with a novel or a familiar conspecific. Both genetic and pharmacological blockage of CB1 receptor in mice caused a significant increase in the discrimination indices indicating that CB1 receptor have a role in social memory (Litvin et al., 2013). Improved performance of CB1 knockout mice in an active avoidance task has been reported (Martin et al., 2002). Blockage of the CB1 receptor in adult mice is associated with improvement in working memory tasks observed in an avoidance task in the elevated T-maze suggesting significant improvement in memory acquisition and consolidation due to the suppression of the CB1 receptors (Takahashi et al., 2005). On the other hand, activation of CB1 by an agonist either by systemic administration or by direct injection into the hippocampus induced memory impairments in a water maze task and in a T-maze (Cha et al., 2006; Abush and Akirav 2010). Activation of CB1 receptors in the medial-prefrontal cortex via adeno-associated virus vector-mediated gene transfer induced impairment in the recognition memory (Klugmann et al., 2011).

The negative effects of CB1 activation on learning and memory could be due to changes in the level of anxiety mediated by CB1 activation because anxiety facilitates learning and memory (Litvin et al., 2013). Anxiety is a type of emotional response that involves the release of diverse neurotransmitters from different brain regions. CB1 receptors control anxiety because of their retrograde function that inhibits the release of neurotransmitters (Ruehle et al., 2012). Genetic or pharmacological manipulation of the CB1 receptors is associated with prominent effects on anxiety and the high density of expression of CB1 receptors in the brain areas, such as the amygdala that mediates emotions, strengthens the concept of the involvement of CB1 receptors in anxiety behavior (Kathuria et al., 2002). CB1 activation has a biphasic effect on anxiety as demonstrated by low doses of agonists inducing an anxiolytic response whereas high doses of agonists induce an anxiogenic response (Vivors et al., 2005). In addition, CB1 receptors in different brain regions differentially regulate anxiety. In the medial prefrontal cortex and dorsal hippocampus (Rubino et al., 2008; Libosa et al., 2015), CB1 activation induced by administration of either direct or indirect agonists produced anxiolytic effects in the elevated plus maze. Whereas, administration of direct agonist in the amygdala produced anxiogenic effects (Rubino et al., 2008). Anxiogenic effects were also observed following systemic administration of indirect agonists (Manduce et al., 2015). In contrast, Libosa and colleagues (2015) reported anxiolytic effects following CB1 activation by an indirect agonist in the amygdala. The basis for this disparity may be due to the fact that anxiety behavior is sensitive to the state of emotionality of the animal where anxiolytic behavior is displayed when tested under low emotional arousal but a state of high emotional arousal masks the anxiety (Morena et al., 2016).



Resident intruder tests and chronic unpredictable mild stress models (CMS) are two valid animal behavioral paradigms used to measure aggression and depression, respectively. To investigate the role of the CB1 receptors in emotional responses, CB1 knockout mice were tested in the resident intruder test and the CMS. Anhedonia, a loss of responsiveness to a pleasurable event, is a measure of depressive behavior exhibited by animals during CMS. The number of attacks made by an intruder when it is placed into a resident's cage measures aggressive behavior. CB1 knockout mice exhibited decreased preference for sucrose water in the CMS and increased intruder attacks in the resident intruder test (Martin et al., 2002).

The CB1 receptors are also associated with social behaviors that are an integral part of learning and developing skills necessary for survival of a group or species. Several studies have addressed the role of CB1 receptors on social behaviors by examining the effects of CB1 agonists or antagonists on play behavior, the earliest form of social behavior (Trezza et al., 2010). CB1 activation by systemic administration of either direct or indirect agonists was found to be associated with a reduction in play as demonstrated by decreased frequency of play solicitation and reciprocation (Trezza and Vanderschuren 2008a, 2008b). In contrast, another study reported an increase in the overall play activity induced by CB1 activation. Pinning is one of the characteristic activities performed by a pair of rats when they are subjected to play. This activity is an important measure of play behavior because execution of pinning happens by equal involvement of both partners. CB1 activation by an indirect agonist induced an increase in the number of pinning in playing rats and this increase was masked in presence of CB1 antagonist (Trezza and Vanderschuren 2009). A similar finding was also reported by Manduce et al. (2015) where CB1 activation by endogenous cannabinoids increased the amount of pinning and pouncing during play

sessions in adolescent rats. Another line of thought concerning the mechanistic basis of social play is the hypothesis that there is a co-activation of the cannabinoid, opioid, and dopamine receptor system during the social play. Social play has been reported to be reduced in rats that were treated with a direct CB1 agonist. However, administration of an indirect agonist enhanced social play and this involved co-activation of the dopamine and opioid receptor systems along with CB1. This enhancement of play induced by the activation of CB1 disappeared in the presence of either an opioid or a dopamine receptor antagonist. In contrast, the presence of a CB1 antagonist can suppress the enhancing effect of the opioid agonist morphine on social play (Review: Trezza et al., 2008).

Considering that CB1 activity is strongly influenced by  $\Delta$ 9-THC exposure and that there is a significant role for CB1 receptors in brain development as well as in mediating anxiety and social behavior, it is possible that exposure to  $\Delta$ 9-THC during a critical window of brain development can have a long-term impact on these types of behaviors. This research will address this issue by testing the following central hypothesis: **Repeated exposure of juveniles to  $\Delta$ 9-THC alters the CB1 signaling pathways leading to long-term effects on behaviors related to anxiety and social interactions.**

### **1.3 Review of Literature**

#### **1.3.1 Elevated plus maze (EPM)**

The EPM is a widely accepted and validated animal model designed to measure the anxiety level in rats and mice (Hogg 1996). The EPM apparatus consists of two closed arms and two open arms of equal size and is elevated to a considerable height above the ground. The open arms have a higher illumination compare to the closed arms. The anxiety level in rats

or mice is predicated based on their natural tendency to avoid illuminated areas. Both forced and voluntary passages into the open arm are associated with increased plasma corticosterone levels and increased freezing which are hormonal and behavioral changes indicative of increased anxiety, respectively (Pellow et al., 1985). The natural exploratory behavior of rat and mice normally occurs more in the closed arms and the time spent in the closed arms increases when the animals are exposed to anxiogenic compounds. However, administration of anxiolytic compounds reduces the natural aversion to the open arms and increase the exploration and time spent in the open arms (Hogg 1996). In recent studies, the EPM has also been used to measure locomotor activity which is indexed by the exploration within the closed arm (Brujinzeel et al., 2016) and the number of closed arm entries (Huang et al., 2010). Table 1.1, below, indicates different types of behaviors reflected by exploration of animal in different parts of EPM.

Table 1.1 Specific behaviors represented by level of exploration performed by rats or mice in different arms of elevated plus maze

EMP Parameters	Associated Behavior	Reference
Fraction of time spent in open arm exploration	Anxiety behavior	Hogg 1996
Fraction of time spent in closed arm exploration	Anxiety behavior Locomotor activity	Hogg 1996 Brujinzeel et al., 2016

Table 1.1 (continued)

Fraction of number of open arm entries	Anxiety behavior	Komada et al., 2008
Number of closed arm entries	Locomotor activity	Huang et al., 2010

There are a fair number of studies examining the effect of  $\Delta 9$ -THC on anxiety by using the elevated plus maze. One study reported that administration of  $\Delta 9$ -THC at a dosage of 0.75 mg/kg (i.p) 30 min before the trial in adolescent male rats can significantly induce anxiolytic effects as demonstrated by increased open arm exploration. This study further examined the potential involvement of serotonin receptors (5HT1A) in the  $\Delta 9$ -THC-induced anxiolytic effects and found attenuation of  $\Delta 9$ -THC-induced anxiolytic effects in presence of a 5HT1A antagonist (Braida et al., 2007). Similar findings were also reported by another study which observed anxiolytic behavior in adolescent rats receiving  $\Delta 9$ -THC (i.p) in the dosage range of 0.075-1.5 mg/kg 30 min prior to testing in the EPM (Rubino et al., 2007). This study suggested the mechanistic basis of CB1-mediated anxiety involves the activity of several kinases.

cFOS is an activator protein that forms transcriptional factor by dimerization with members of jun family and can be used as a mapping tool to resolve the neuronal circuits underlying the behaviors induced by stress including anxiety (Kovacs 1998). CREB is a transcriptional factor that is activated in response to cAMP and associated with depressive or emotional behaviors (Barrot et al., 2002). CREB-mediated gene expression is closely

related to the activity of kinases such as protein kinase A (PKA) which phosphorylates CREB and calcium/calmodulin protein kinase II (CaMKII) which inhibits CREB by preventing dimerization and binding of CREB-related proteins (Rubino et al., 2007). In the EPM, cFOS phosphorylation was increased in behavioral control rats compared to rats that did not go through testing.  $\Delta$ 9-THC treatment altered this scenario in that there was a decrease in the level of phosphorylated cFOS in the prefrontal cortex (PFC) and amygdala. In the hippocampus, testing in the EPM did not have any effect on CREB, CaMKII, PKA, and extracellular-signal-regulated kinase (ERK) signaling.  $\Delta$ 9-THC treatment also altered this scenario by reducing the activity of CaMKII which, in turn, triggers CREB activation. All of the  $\Delta$ 9-THC-induced effects observed in this study were reversed in the presence of the CB1 antagonist AM251 which suggests CB1-mediated effects on all of these kinases and transcription factors. In PFC, there was an inhibitory effect of EPM on CaMKII, but there was no prominent effect of  $\Delta$ 9-THC (Rubino et al., 2007). This disparity of effects in different regions could be due the fact that these three regions differentially modulate the anxiety induced by  $\Delta$ 9-THC. Direct administration of  $\Delta$ 9-THC into the hippocampus and PFC produced anxiolytic effects whereas, direct administration of  $\Delta$ 9-THC into the amygdala led to anxiogenic effects (Rubino et al., 2008b). On the other hand, an anxiogenic response was observed at the dosage of 2.5 mg/kg  $\Delta$ 9-THC (i.p) given to adolescent and adult rats 30 min before the EPM trial with adults showing significantly more effects at the same dosage (Schramm-Sapyta et al., 2007). An anxiogenic response was also observed in adolescent rats who were given gradually increasing doses of  $\Delta$ 9-THC on the scale of 2.5-10 mg/kg for 11 consecutive days. This anxiogenic response persisted for seven days following the last exposure (Stopponi et al., 2014). However, no effects were observed 30

days following the administration using the same dosing scale and regimen (Rubino et al., 2008a).

The most common route of exposure to  $\Delta$ 9-THC in humans at adolescent ages is through inhalation in the form of smoke by  $\Delta$ 9-THC cigarettes or pipes. Recently, Brujinzeel and colleagues (2016) simulated this exposure paradigm by exposing 55 days old male rats to cannabis smoke containing 5.7%  $\Delta$ 9-THC. The concentration of  $\Delta$ 9-THC in the serum following the exposure was found to be 225 ng/ml which is similar to the level detected in the humans following smoking cannabis. This study observed no significant effects on open arm activity but did observe an increase in exploration of closed arm in treated rats as compared to control rats. This suggests that  $\Delta$ 9-THC-induced increase in locomotor activity which was then confirmed by using open field test. This increased locomotion was only observed when tested immediately after the exposure and the effects were no longer detected at 4 hours following exposure. However, this time-line of appearance and disappearance of effect does not always hold true because other studies reported contradictory findings. Chronic administration of 3 mg/kg (i.p)  $\Delta$ 9-THC to male mice at adolescent or adult ages for three weeks did not change the level of anxiety in any group when the EPM test was performed immediately following treatment. When the EPM test was performed six weeks following the cessation of exposure, there was an increased aversion to the open arms in the treated rats (Murphy et al., 2017). Another study investigated the effect of 10 mg/kg  $\Delta$ 9-THC administered subcutaneously in male mice daily for 10 consecutive days then followed by administration of CB1 antagonist SRA141716 (i.p) on the last day in sub-samples of both  $\Delta$ 9-THC and vehicle mice. Mice that received  $\Delta$ 9-THC alone performed in the same way as controls in terms of percent

open arm entries and percent time in the open arms.  $\Delta 9$ -THC-treated mice that were given a challenge of 3 mg/kg of SRA141716 exhibited a sharp decrease in the percent time in the open arms and percent open arm entries as compared to their respective controls. These data suggest the anxiety inducing effect of  $\Delta 9$ -THC withdrawal (Huang et al., 2010), which is also observed in humans (Winstock and Lea 2009). However, rats and mice may have a different time-line of anxiogenic effects following  $\Delta 9$ -THC withdrawal because Rubino and colleagues (2008a) observed no effects in rats 30 days after the last exposure but Murphy and colleagues (2017) observed effects in mice following 42 days after the last exposure (Murphy et al., 2017).

### **1.3.2 Social play**

Play is a deceptive term as it not so easily defined as it sounds because of the ambiguity in the categorizing the behavior as “playful behavior”. Play is associated with many aspects of development and especially important is the fact that it provides a supportive environment in which children can develop the skills to interact with others. In most mammals, play is generally observed between the ages of weaning and sexual maturation. This phenomenon is very important in juvenile animals and one of the determining factors in establishing adult behavior (Anne and Eimon, 1981). There are many theories concerning the concept of play behavior but the empirical evidence to actually define the boundaries of actions involved in play are still questionable. This may be because actions involved in play are so random and seem to be performed without any purpose (Panksepp et al., 1984). Because of the ambiguity in these actions and the absence of a specific purpose in performing them, the biochemical mechanism that regulate the programming of this behavior is not well characterized (Blake and McCoy, 2015). However, play is thought to

be tightly regulated by some unidentified brain circuits not only in the short term but throughout the life span of an organism (Panksepp et al., 1984).

### **1.3.2.1 Social play in developmental disorders**

It has also been determined in human juveniles that play behavior is necessary for the development of normal social interactions. Play behavior therapy is currently being used as a treatment in several neurodegenerative disorders (Dawson 2008). Social play has a prominent role in identification and diagnosis of the autism spectrum disorder. Autism is a developmental disorder characterized by dysfunctions in specific aspects of behavior including the ability to participate in social interactions, the ability to communicate, and characterized by repetitive behaviors and lack of interests (Hille 2004). These deficits are similar to alterations in social play as evidenced by the significant differences in social play activities observed in children diagnosed with autism. In addition, a modified form of play that would facilitate autistic children to engage in play could be an effective therapy in autism (Restall and Magill-Evan 1993). Alterations in play behavior is also associated with Attention Deficit Hyperactivity Disorder (ADHD). When observing patterns of play, pre-school children with ADHD exhibited different patterns of play as compared to non-ADHD children. In ADHD children, play activities were decreased while non-play activities were increased. There was also a decrease in the sustained attention of these children and an increase in the frequency of transition from one activity to another (Alessandri 1992).

### **1.3.2.2 Structure and neurobiology of play**

Based on our present knowledge about most behaviors, the molecular mechanisms controlling a specific behavior are not well characterized and there is no single gene



directly involved in the control of certain behaviors (Peills and Peills 1993, Popova 2006, Blake and McCoy 2015). This is especially true for play. Play is a complex form of behavior involving a variety of random actions performed by the individuals involved that are not only difficult to capture in words but the boundaries between the different forms of these random actions are even difficult to define. Despite play being a very complex and random phenomenon, it does have a distinct role in the establishment of normal social, sexual, and aggressive behavior in adults and it is one of those essential tasks required for the normal development of the brain (Panksepp et al., 1984).

In an experimental setup, social play in rats can be observed by documenting the occurrence of certain parameters such as pouncing, chasing, grooming wrestling, napping, pinning, sniffing and crawl over. A bout of play generally starts by one rat approaching or soliciting another to play. The soliciting rat attempts to sniff or rub the neck of the play partner and from this, chasing, boxing, wrestling crawling, and/or pinning may follow. The time that two rats spend involved in these activities is considered as the time of play. Each of these activities has been validated to represent specific type of behavior (Table 1.2). From the neurobiological point of view, social play is influenced by multiple centrally acting neurotransmitters such as acetylcholine, adenosine, catecholamine, dopamine, serotonin, opioids, and noradrenaline (Vanderchuren et al., 1997).

Table 1.2 Different types of activities performed during the act of play and associated behaviors represented by each activity

Play Activities	Behavior	Reference
Sniffing/anogenital sniffing	Contact behavior	Peills and McKenna 1995.
Self-Grooming	Aggressive behavior	Miczek 1974
Crawling over/under	Unfamiliar situation	Peills and McKenna 1995.
Chase	Appetitive play	Panksepp and Beatty 1980.
Nape attack	Play solicitation	Peills and McKenna 1995.
Pinning	Play variable	Panksepp and Beatty 1980.
Wrestling/Time of wrestling	Consummatory play	Panksepp and Beatty 1980.

### 1.3.2.3 Effect of $\Delta 9$ -THC on play behavior

Miczek (1974) first reported the effect of  $\Delta 9$ -THC on aggressive interactions by observing a dominant adult male rat and a subordinate adult male rat inside a dark chamber. When the subordinate rats were administered 4 mg/kg (i.p) of  $\Delta 9$ -THC, the defensive behaviors of the submissive rats were decreased as evidenced by the increased incidence of severe injuries inflicted by the dominant rat during the test. When the dominant rat was administered the same dosages of  $\Delta 9$ -THC, there was reduction in the frequency of aggressive attacks towards the subordinate conspecific. In a subsequent study, injection of 5 mg/kg of  $\Delta 9$ -THC (i.p) into adult mice resulted in a reduction in exploratory behavior and an increase in immobility in male-to-male encounters. In male-female encounters,

there was a reduction in sexual behaviors as indicated by decrease in the number of mounts and chases followed by sniffs (Cutler and Mackintosh 1984).

Active social interaction of two rats placed into the same arena can be measured by the time it takes for the rats to approach each other or for one rat to move towards a stationary conspecific. Higher latency indicates a reduced level of interaction. As reported by Malone and colleagues (2009), the latency of initial interaction was reduced in adolescent rats treated with 1 mg/kg of  $\Delta$ 9-THC (ip) but this reduction was not observed at higher dosages (3 or 10 mg/kg). Similarly, an increase in investigative behavior was observed in adult rats (PND90) receiving 2 mg/kg of  $\Delta$ 9-THC (sc) during PND2-10 (Newsome and Kelley, 2008). However, during exposure, no effect on total social interactions were observed in adult and adolescent rats receiving a chronic dosages of 5 mg/kg  $\Delta$ 9-THC (ip). However, when a social interaction test was performed 16 days following the last administration of  $\Delta$ 9-THC, there was a decline in social interactions in the treated rats (Quinn et al., 2008). A decline in social interactions as evidenced by reduced body sniffing and allo-grooming and increased rearing was observed in adult rats receiving 2 mg/kg of  $\Delta$ 9-THC as compared to saline-control rats. Interestingly, this study also reported no differences in social interactions between  $\Delta$ 9-THC-treated rats and ethanol-control rats (Slamberova et al., 2016) suggesting that it is possible that ethanol, which is the most frequently used solvent in  $\Delta$ 9-THC studies, may have some effect on behavioral performance in rats.

#### **1.4 Proteomic Approach to Study Behavior**

Proteomics involves the study of complete protein complement of the genome of an organism. Recently, this technology has been employed to study the protein profiles in

rodent brain to study behavioral and/or developmental changes (Richard et al., 2002). The gel-based proteomics is one of the classic methods used for protein identification and quantification. This method involves the separation of proteins by using two dimensional polyacrylamide gel electrophoresis (2D PAGE) and identification of proteins by using Mass Spectrometry (MS) or Tandem Mass Spectrometry (MS/MS; Chevalier 2010). Gel-based proteomics has been employed to study the changes in the protein expression in mice with anxiety like behavior and psychomotor deficits. The differentially expressed proteins were then analyzed in Ingenuity Pathway Analysis (IPA; Qiagen Inc., <https://www.qiagenbioinformatics.com>) to predict the biofunctional categories in which these differentially expressed proteins were involved in (Abdullah et al., 2011). Similarly, the gel-based proteomic technique was used to evaluate the differentially expressed proteins in developing brains of rats and mice (Quinn et al., 2008; Taoka et al., 2000; Fountoulakis et al., 1999) or to quantify the differentially expressed proteins in different brain regions obtained from human Alzheimer's Disease patients (Schonberger et al., 2001). The gel-based proteomics method can also be modified to specifically identify oxidatively modified proteins or phosphorylated proteins in tissues to obtain more specific information about experimental or disease conditions (Butterfield et al., 2006). Another method utilized in proteomics is the label free proteomic approach, which involves digestion of individual protein samples followed by separation of digested peptides by using liquid chromatography. The separated peptides are then introduced into a mass spectrometer to generate the MS/MS spectra for each identified peptide (Zhu et al., 2010). The number of spectra matched to specific peptide in a protein is a linear measure for relative abundance of the protein (Zhang et al., 2006). In this study, we employed the label free shotgun

proteomic approach to study the expression pattern of proteins in the amygdala of adolescent rats exposed to  $\Delta 9$ -THC as juveniles.

Considering the increased availability of marijuana edibles and the adverse effects of its active component ( $\Delta 9$ -THC) on brain development, this research was designed to investigate the long-lasting effects of  $\Delta 9$ -THC ingestion at early stage of life in rat model. Chapter II will investigate the  $\Delta 9$ -THC-mediated effects on anxiety level and performance during social interaction test at adolescent stage. Chapter III will not study behavior but will address proteins and canonical pathways affected following repeated  $\Delta 9$ -THC ingestion. Chapter IV will analyze the canonical pathways involved in the social behaviors, and alterations in these canonical pathways following repeated  $\Delta 9$ -THC ingestion. Chapter V will be the overall conclusion emphasizing the adverse behavioral outcomes, and alterations in canonical pathways at adolescent ages due to unwanted activation of CB1 receptors at juvenile ages.

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CHAPTER II  
EFFECT OF REPEATED JUVENILE EXPOSURE TO  
 $\Delta$ 9-TETRAHYDROCANNABINOL ON ANXIETY-RELATED BEHAVIORS AND  
SOCIAL INTERACTIONS IN ADOLESCENT RATS

**2.1 Introduction**

Cannabinoids are a group of terpenophenols present in the granular trichomes of the plant *Cannabis sativa*. Because of the psychoactive properties, cannabis preparations, such as marijuana and hashish, have been used recreationally for centuries and remain one of the most widely used illicit drugs (Bruijzeel et al., 2016; Substance Abuse and Mental Health Services Administration, 2014; Fellermeier et al., 2001). After the discovery of the potential medical application of marijuana, there has been much debate about its legalization for medical use. The debate was initiated in 1972 when the National Organization for the Reform of the Marijuana Laws (NORML) filed a petition with the Bureau of Narcotics and Dangerous Drugs to reschedule marijuana under the Controlled Substances Act. However, crude marijuana remains classified with the Schedule I drugs, which are described as ‘drugs with no accepted medical applications in treatment in United States’ (Zeese, 1999). Despite the strict regulation of marijuana, its use has increased dramatically in the past few decades. This widespread use is a growing concern

especially because of the abuse of the drug by pregnant women and by adolescents. In fact, there are many behavioral and neurological disorders in human and animal studies associated with marijuana exposure either during the prenatal period (Higuera-Mates et al., 2015) or during adolescent ages (Fontes et al., 2011; Fried et al., 2002; Meier et al., 2012; Pope et al., 2001). However, exposure to marijuana is also increasing in a very unlikely population. From 2006 through 2013, the rate of marijuana exposure among children 5 years old or less rose 147.5% in the United States. In states that have legalized marijuana, the exposure rate increased almost 610% during the same period (Onders et al., 2016). An even earlier study reported that marijuana exposure among children 9 years old or less increased by 30.3% per year between 2005 and 2011 in decriminalized states but did not change in nonlegal states (Wang et al., 2013). In both studies, these data were obtained from reports to Poison Control Centers and it is therefore likely that many additional exposures are occurring that go unreported. While literature has expanded in recent years with respect to the behavioral and neuroanatomical effects of adolescent and prenatal exposure to marijuana in laboratory animals, very little information is available on the persistent effects of marijuana exposure during developmental periods that mimic exposure in young children.

The main psychoactive component of marijuana is  $\Delta^9$ -tetrahydrocannabinol ( $\Delta^9$ -THC) (Gaoni & Mechoulam 1964) which activates the cannabinoid receptors, including type I cannabinoid receptor (CB1). CB1 receptors are abundant in the brain (Devane et al., 1988), appear during fetal development (Berrendo et al., 1998), and the highest density of CB1 receptors occur in brain regions which are directly involved in brain developmental processes (Fernandez-Ruiz et al., 2000). During development, the CB1

receptor is involved in synaptogenesis, the formation of connections across the neurons which is essential for efficient processing of adult cognition (Basavarajappa et al., 2009). Synaptogenesis peaks during the postnatal period suggesting that any inappropriate activation of the CB1 receptor during this postnatal period, such as that which occurs during  $\Delta 9$ -THC exposure, could result in altered synaptic connections. This could lead to neurobehavioral abnormalities that may not be observed until later stages of life. In fact, many behavioral abnormalities have resulted from developmental  $\Delta 9$ -THC exposure in animal studies. Most of these studies have focused on the behavioral abnormalities observed following  $\Delta 9$ -THC exposure either during pregnancy (Hutchings et al., 1991; Newsome & Kelly 2008; Rubio et al., 1995, 1998) or adolescence (Fleming et al., 2016; Keeley et al., 2015; Zamberletti et al., 2012; Realini et al., 2011; Rubino et al., 2011; Quinn et al., 2008; Rubino et al., 2008; Cha et al., 2006; Jarbe et al., 1998; Miczek & Barry 1974; Ueki et al., 1972).

Since the unintentional exposure of marijuana in children is increasing dramatically (Onders et al., 2016; Wang et al., 2013), it is imperative that research focus on investigating whether or not this pediatric exposure scenario can lead to persistent abnormalities. The majority of pediatric exposures result from the ingestion of  $\Delta 9$ -THC-infused edibles which are indistinguishable from their equivalent non-infused food product with respect to look, smell, and taste (Wang et al., 2017). For commercial products, several states with legalized marijuana (Alaska, Oregon, Colorado, and Washington, California) have limited the amount of  $\Delta 9$ -THC that can be in a single serving (5-10 mg; Gourdet et al., 2017). However, multiple servings of  $\Delta 9$ -THC are allowed in a single edible (i.e., brownie, cookie, chocolate) and children could easily

consume multiple servings and even multiple edibles (Gosh et al., 2015). Even with limitations placed on the number of servings, the total  $\Delta 9$ -THC concentration in one edible can reach as high as 100 mg. Some individual edibles contain up to 1000 mg (Richards et al., 2017). In addition, the exposure levels from homemade  $\Delta 9$ -THC-infused edibles, which are not restrained by regulations, could be even higher. Thus, the range of exposure levels in children can be very wide. For this investigatory project, we selected 10 mg/kg/day which has been utilized in previous studies exposing developing rodents (Burston et al., 2010; Moore et al., 2010; Philippot et al., 2016). The human equivalent dose for this dosage in a 5 year old child (17-20 kg) calculates to be 40-48 mg/day. However, it is not clear if this dosage recapitulates exposure levels in children. This research project was designed to determine if exposure to the psychoactive component of marijuana  $\Delta 9$ -tetrahydrocannabinol during the postnatal period altered anxiety and social behaviors in adolescent rats. Since rat brain development at postnatal day (PND) 7 has historically been used as a rough estimate of human brain development at birth (Clancy et al., 2007), the exposure treatment period utilized in this project was from PND10-16 which should be equivalent to early childhood ages in humans (Beck et al., 2011).

## **2.2 Materials and Methods**

### **2.2.1 Chemicals**

$\Delta 9$ -tetrahydrocannabinol was provided by the National Institute on Drug Abuse. All other chemicals were purchased from Sigma Chemical Co. (St. Louis, MO).

### **2.2.2 Animals**

A breeding colony of adult male and female Sprague Dawley rats (CD IGS; Harlan Laboratories, Indianapolis, IN) were housed in an Association for Assessment and Accreditation of Laboratory Animal Care-accredited facility with temperature control ( $22 \pm 2^\circ\text{C}$ ) and with lights on between 0700 and 1900 for a 12-h dark-light cycle. Tap water and LabDiet rodent chow were freely available during the experimentation. The procedures used in this project were approved by the Mississippi State University Institutional Animal Care and Use Committee. From this colony, male and female rat pups were obtained and the day of birth was considered as PND0. On PND10, rat pups within the same litter were assigned to different treatment groups. There were always a representative control animal of the same sex present in each litter to match the  $\Delta 9$ -THC-treated animal to ensure equal distribution of animals to different treatment groups.

### **2.2.3 Exposure**

Beginning on PND10, rats were exposed daily for 7 days to either corn oil (vehicle) or 10 mg/kg  $\Delta 9$ -THC by oral gavage at a volume of 0.5 ml/kg. Solutions were delivered to the back of the throat using a 50- $\mu\text{l}$  tuberculin syringe equipped with a 1-inch 24-gauge straight intubation needle (Popper and Sons, Inc., New Hyde Park, NY). At weaning, the rats were weaned and remained housed with littermates throughout behavioral testing.

### **2.2.4 Elevated plus maze (EPM)**

The elevated plus maze was constructed of black-painted wood and consisted of four arms (50 X 10 cm), two open and two enclosed (40 cm wall), intersecting at perpendicular distances to each other to form a  $90^\circ$  angle and elevated to a height of 50

cm above the floor. On PND29, the rats were transported to the testing room, placed in the center of the maze and allowed to explore the maze for 5 min. After each test, the maze was cleaned with 70% ethanol and dried. Each test session was recorded using a remotely operated Canon EOS Rebel digital camera. The videos were scored by two observers who were blind to treatment.

The observers scored the following behavioral parameters: (1) number of entries into the open arms, (2) number of entries into the closed arms, (3) time spent in the open arms, (4) time spent in the closed arms, and (5) time spent in the central square. From these data, the (6) percent open arm entries and (7) percent open arm time were calculated. In addition, the following ethological oriented parameters were scored: (8) number of stretch-attend postures (defined as exploration of the open arm with the front part of the body, while the hind region remains in the closed arm) and (9) number of head-dips (protruding the head over the ledge of an open arm and downwards towards the floor regardless of the location of the animals body). Entries into closed arm or open arm were counted only when all four paws including tail of rats were in respective arms.

Different anxiety levels have been observed under different levels of illumination following stimulation of the endocannabinoid system in rats (Haller et al., 2002). Thus, testing was conducted under two different illumination scenarios. The first was high illumination in which the illumination was ~600 LUX in the open arms and ~100 LUX in the closed arms. For this scenario, the controls (n=11 of each sex) were derived from 8 litters and the  $\Delta$ 9-THC treated (n=12 of each sex) were derived from 10 litters. The second was low illumination in which the illumination was ~10 LUX in the open arms and ~2 LUX in the closed arms. For this scenario, the controls (n=26-30 of each sex)



were derived from 22-23 litters and the  $\Delta$ 9-THC treated (n=20 of each sex) were derived from 15-16 litters. There were never more than two animals of the same treatment and sex from same litter used in any aspect of this study.

### **2.2.5 Social behavior (PND37-38)**

Since previous behavioral test can affect the performance in subsequent behavioral test (Blokland et al., 2012; Mellwain et al., 2001), social behavior was conducted only on rats tested under low illumination in the elevated plus maze. The low illumination was selected over high illumination because, intuitively, it was considered to be less invasive and exerts less stressful effect. The social play test requires two rats of the same treatment in order to obtain a single replication and this is the reason for there being a greater numbers of litters and rats tested under low illumination in the elevated plus maze as compared to those tested under high illumination. The behavioral arena was a clear empty litter-cage with bright light (600 lux). Following a 24 hour isolation period, two rats of the same treatment, sex, age and size but from different litters were placed into different corners of the cage and allowed to interact for 10 min. Each test session was recorded using a remotely operated Canon EOS Rebel digital camera. After each test, the cage was emptied, cleaned with 70% ethanol, dried, and refilled with fresh litter. The videos were scored by two observers who were blind to treatment.

The observers scored the behavioral parameters as previously described (Ku et al., 2016; Himmler et al., 2013; Trezza et al., 2010; Varlinskaya & Spear 2002; Vanderschuren et al., 1997). These included: (1) time to first interaction, (2) frequency of self-grooming, (3) frequency of social exploration (body or anogenital sniffing or licking), (4) frequency of crawling over and under, (5) frequency of chasing, (6) frequency of pouncing/nape

attacks (playful attacks directed at the nape of the neck), (7) frequency of play fighting (boxing and wrestling), (8) frequency of pinning (one animal lying with its back on the floor with the other animal standing over it), and (9) the time spent play fighting. When scoring these parameters, the behavior of each individual rat was not recorded and the score obtained is a combined measure of the occurrence of the behavior in both rats regardless of which rat performed/initiated the behavior.

### **2.2.6 Statistical analysis**

All data were subjected to testing by analysis of variance (ANOVA) using the SAS statistical package (SAS Institute Inc., Cary, NC). For body weights, the data were tested using the general linear model with a repeated measures paradigm and was found to violate the assumption of sphericity. Therefore, subsequent analysis using the Mixed procedure (Littell et al., 1996) was conducted with a repeat measures paradigm with a Huynh-Feldt covariance structure (Huynh & Feldt 1970), followed by separation of means using least squares mean (LSM). The analysis identified significant differences in the main effects (sex, treatment, and day) and all possible interactions.

Behavioral data were analyzed by ANOVA using the Mixed procedure followed by mean separation by Least Square Means (LSM). Prior to analysis, the distribution of the conditional residuals for the behavioral data were evaluated for each parameter to determine the appropriateness of the statistical model. Any parameter not found to be sufficiently normally distributed were subjected to Blom's transformation in SAS using the PROC RANK procedure with the NORMAL option (Altman 1991). The analysis identified significant differences in the main variables (sex, treatment and day) and all possible interactions. If either main effect was significant or tending towards

significance, an additional lower level analysis (one-way ANOVA) was performed to determine differences between sexes within each treatment or treatment within each sex. The criterion for significance was set at  $p < 0.05$ .

## **2.3 Results**

### **2.3.1 Weight gain**

Oral exposure to 10 mg/kg  $\Delta 9$ -THC significantly reduced the amount of weight gained in both females and male pups (Figure 1) with no statistically significant difference between sexes. No signs of toxicity were observed in the pups. The pups were not lethargic and did not exhibit any abnormal movements or behavioral patterns during the exposure period. By the time of behavioral testing, there were no longer any statistically significant differences between treatment in either sex. However, male rats weighed significantly more than female rats ( $P=0.0001$ ).

### **2.3.2 Elevated plus maze**

For the elevated plus maze, the parameters measured under high illumination and the parameters measured under low illumination were analyzed separately. For both high and low illumination, the overall statistical analyses indicated that there was no significant overall effect of treatment and no significant treatment  $\times$  sex interaction on any activity parameter associated with any open arm activity but some significant differences were detected in certain parameters. When detected, an additional lower level analysis was conducted to determine the presence of either sex differences within each treatment or treatment differences with each sex.

### 2.3.3 Open arm activity

There was no significant effect of treatment and no significant treatment  $\times$  sex interaction in any parameter associated with open arm activity tested under either high or low illumination. However, statistically significant sex differences were detected in certain parameters.

There was a significant overall effect of sex in the number of entries into the open arm under both high illumination ( $F_{(1,32)} = 7.8$ ,  $p = 0.089$ ) and low illumination ( $F_{(1,46)} = 7.57$ ,  $p = 0.0085$ ). Under high illumination, males made significantly more open arm entries than females (Figure 2A). The effects appear to be more significant between males and females in the  $\Delta 9$ -THC treated group (post hoc analysis,  $p = 0.0145$ ) with no significant difference between sexes in the control group. Under low illumination, the pattern was reversed with females making significantly more open arm entries than males (Figure 2B) but post hoc analysis did not reveal any treatment related effects. When calculated in terms of % of open arm entries, there was a significant effect of sex only under high illumination ( $F_{(1,32)} = 14.56$ ,  $p = 0.0006$ ). Under high illumination, both  $\Delta 9$ -THC treated males (post hoc analysis,  $p = 0.0175$ ) and control males (post hoc analysis,  $p = 0.0273$ ) had a higher % of open arm entries (Figure 2C) than their respective female treatment group. Under low illumination, females had a slightly higher % of open arm entries than males but there was only a trend towards significance ( $F_{(1,46)} = 3.53$ ,  $p = 0.0668$ ) (Figure 2D).

There was a significant effect of sex in the time spent in the open arms under both high ( $F_{(1,32)} = 5.52$ ,  $p = 0.0253$ ) and low illumination ( $F_{(1,46)} = 4.97$ ,  $p = 0.0308$ ). Under high illumination, males spent significantly more time in the open arms than females (Figure

1E). The effects appear to be more significant between males and females in the  $\Delta 9$ -THC treated group (post hoc analysis,  $p = 0.0386$ ) with no significant difference between sexes in the control group. Under low illumination, the pattern was reversed with females making significantly more open arm entries than males (Figure 2F) but lower level analysis did not reveal any treatment related effects. When calculated in terms of % of time spent in the open arms, there was a significant effect of sex under both high ( $F_{(1,32)} = 5.52, p = 0.0253$ ) and low illumination ( $F_{(1,46)} = 4.97, p = 0.0308$ ). Under high illumination, males had a higher % of time spent in the open arm than females (Figure 2G). The effects appear to be more significant between males and females in the  $\Delta 9$ -THC treated group (post hoc analysis,  $p = 0.0386$ ) with no significant difference between sexes in the control group. Under low illumination, females had a higher % of time spent in the open arm than males but lower level analysis did not reveal any treatment related effects (Figure 2H).

#### **2.3.4 Ethological oriented parameters and overall activity**

Under both high and low illumination, there was no significant effect of treatment and no significant treatment  $\times$  sex interaction in any ethological oriented parameter. However, sex differences were detected in certain parameters.

With respect to the numbers of stretch attended postures made under high illumination, the  $\Delta 9$ -THC treated rats of both sexes made more as compared to their respective controls but was not statistically significant (Figure 3A). Under low illumination, there was a significant overall effect of sex ( $F_{(1,46)} = 4.63, p = 0.0367$ ) with males making more than females. However, this effect appears to be more significant in the  $\Delta 9$ -THC

treated groups (post hoc analysis,  $p = 0.0291$ ) with no significant difference between males and females in the control group (Figure 3B).

With respect to the number of head dips made under high illumination, there was a trend toward significance in the overall effect of sex ( $F_{(1,46)} = 3.77$ ,  $p = 0.0593$ ) with males making more than females. Although not statistically significant, the  $\Delta 9$ -THC treated males made more head dips than did the  $\Delta 9$ -THC treated females with similar levels made between male and female controls (Figure 3C). Under low illumination, there was a trend towards significance for the effect of treatment ( $p=0.0590$ ). This effect appeared to be more significant in males with greater number of head dips in the  $\Delta 9$ -THC treated group than controls (post hoc analysis,  $p = 0.0227$ ) with no significant difference observed between treatment groups in females (Figure 3D).

During elevated plus maze testing, entries into the closed arms can be considered an indicator of activity levels (Cruz et al., 1994). Overall analysis indicated that under both high and low illumination, there was no significant effect of sex and no significant treatment  $\times$  sex interaction. However, there was a trend towards a significant effect of treatment under high illumination ( $F_{(1,32)} = 3.35$ ,  $p = 0.0771$ ) and a significant effect of treatment under low illumination ( $F_{(1,46)} = 6.02$ ,  $p = 0.0180$ ). However, the effect appeared to mainly occur in males where  $\Delta 9$ -THC treated males made more entries into the closed arms than did control males under both high (Figure 3E) and low (Figure 3F) illumination (post hoc analysis,  $p=0.0453$  and  $p=0.0210$ , respectively). The number of entries was similar between the two treatment groups in females regardless of illumination level.

### 2.3.5 Social behavior

There were no significant overall effects of treatment or sex and no significant sex  $\times$  treatment interactions in the time to first interaction, the frequency of self-grooming, the frequency of crawling over and under, the frequency of chasing, the frequency of pouncing, or the frequency of pinning (data not shown). However, treatment effects were observed on the frequency of social exploration (body or anogenital sniffing or licking), the frequency of nape attacks, the frequency of play fighting (boxing and wrestling), and the time spent play fighting. In these four parameters, there was no significant overall effect of sex and no significant sex  $\times$  treatment interaction.

With respect to social exploration, there was a significant overall effect of treatment ( $F(1,30) = 7.65, p = 0.008$ ). Episodes of social exploration were significantly decreased in rats exposed to  $\Delta 9$ -THC as compared to that in control rats (Figure 4A). It appeared that the reduction in social exploration occurred equally in both males and females exposed to  $\Delta 9$ -THC (Figure 4A inset). With respect to the frequency of nape attacks, there was a significant overall effect of treatment ( $F(1,30) = 16.93, p = 0.0004$ ). Rats exposed to  $\Delta 9$ -THC made significantly higher number of nape attacks than did control rats (Figure 4B). Although both  $\Delta 9$ -THC-treated females and males exhibited increased episodes of nape attacks, there appeared to be a greater numbers in males (Figure 4B inset). With respect to the frequency of play fighting, there was a significant overall effect of treatment ( $F(1,30) = 6.35, p = 0.0188$ ). Rats exposed to  $\Delta 9$ -THC engaged in more episodes of play fighting than did the control rats (Figure 4C). This increased level of play fighting was present in both  $\Delta 9$ -THC exposed males and females (Figure 4C inset). With respect to the time spent playing, there was an overall significant difference

of treatment ( $F(1,30) = 6.57, p = 0.0171$ ). Rats exposed to  $\Delta 9$ -THC spent a significantly greater amount of time engaged in play fighting than did control rats (Figure 4D). This increased amount of time spent play fighting was observed in both  $\Delta 9$ -THC exposed males and females (Figure 4D inset).

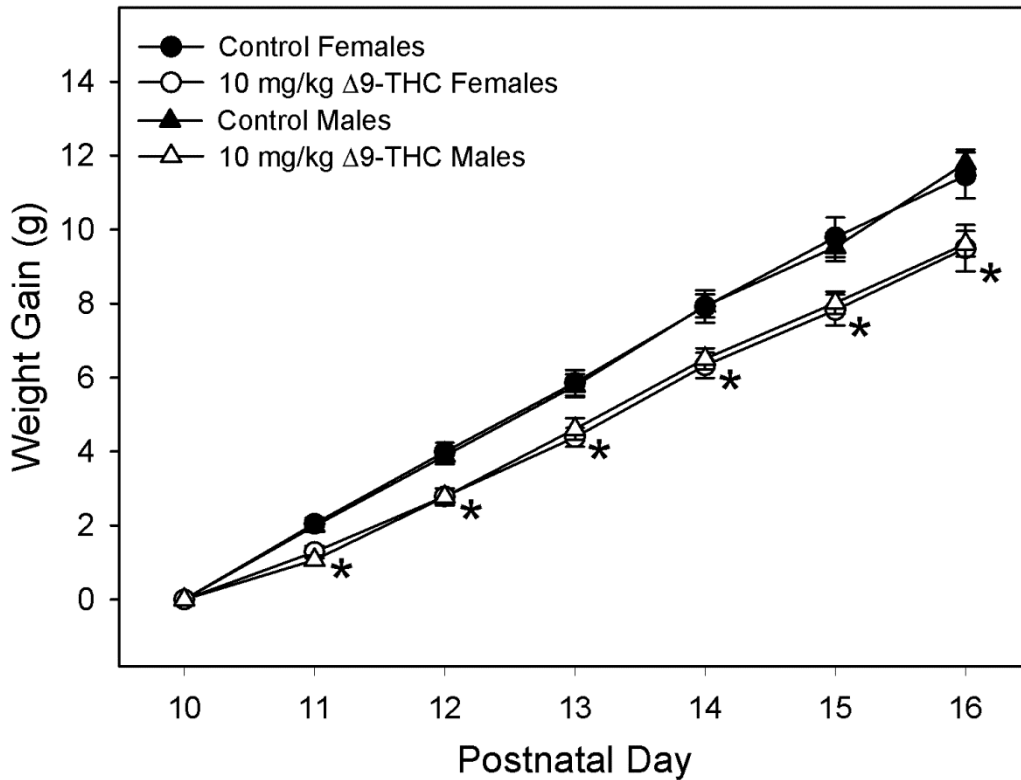


Figure 2.1 Plot of weight gain of male and female rat pups exposed daily to either corn oil (control) or 10 mg/kg of  $\Delta 9$ -THC as juveniles from postnatal days 10-16. Values are expressed as weight  $\pm$  SE ( $n = 31-37$ ). For both sexes, weight gain in treated rats was significantly different from that in the controls ( $p \leq 0.05$ ) at all time points.



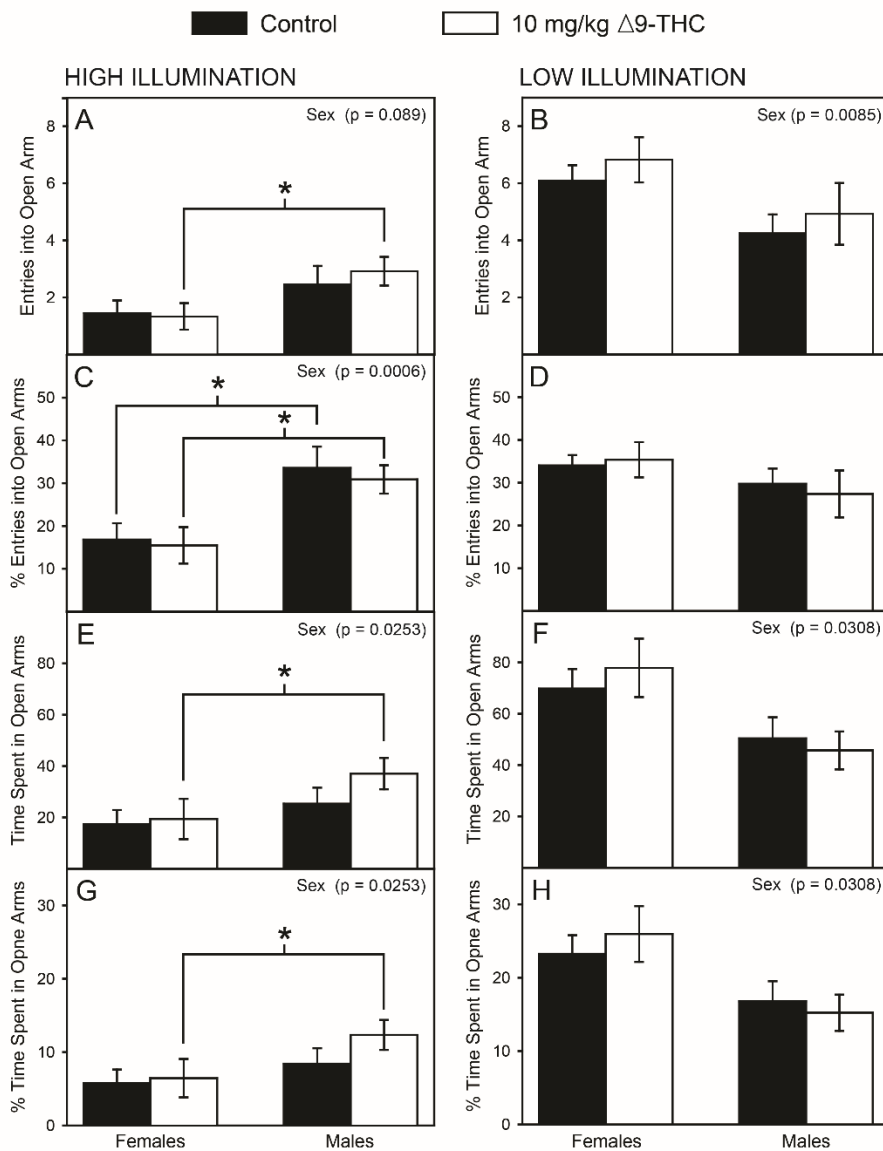


Figure 2.2 Open arm activity in the elevated plus maze under different levels of illumination as described in the Materials and Methods. Number of entries into open arms under high (A) and low (B) illumination, percent entries into open arms under high (C) and low (D) illumination, time spent in the open arms under high (E) and low (F) illumination, and percent time spent in the open arms under high (G) and low (H) illumination, in male and female adolescent rats (PND29) exposed to either corn oil or 10 mg/kg of  $\Delta 9$ -THC as juveniles from postnatal days 10-16. Values are expressed as means  $\pm$  SEM or percent  $\pm$  SEM (n=11-30). Brackets with asterisks (\*) are statistically significant ( $p \leq 0.05$ ) from one another

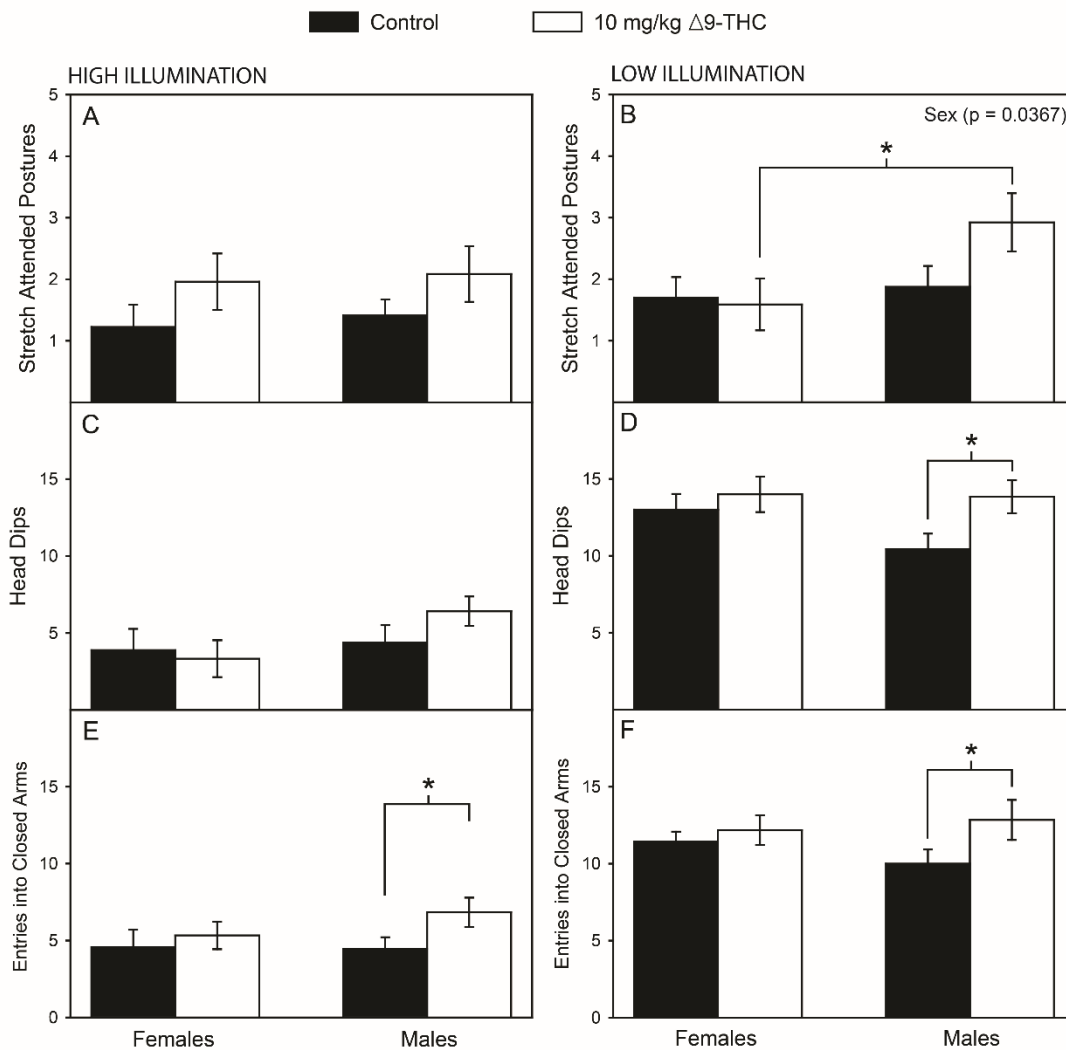


Figure 2.3 Open arm activity in the elevated plus maze under different levels of illumination as described in the Materials and Methods. Number of entries into open arms under high (A) and low (B) illumination, percent entries into open arms under high (C) and low (D) illumination, time spent in the open arms under high (E) and low (F) illumination, and percent time spent in the open arms under high (G) and low (H) illumination, in male and female adolescent rats (PND29) exposed to either corn oil or 10 mg/kg of  $\Delta$ 9-THC as juveniles from postnatal days 10-16. Values are expressed as means  $\pm$  SEM or percent  $\pm$  SEM (n=11-30). Brackets with asterisks (\*) are statistically significant ( $p \leq 0.05$ ) from one another

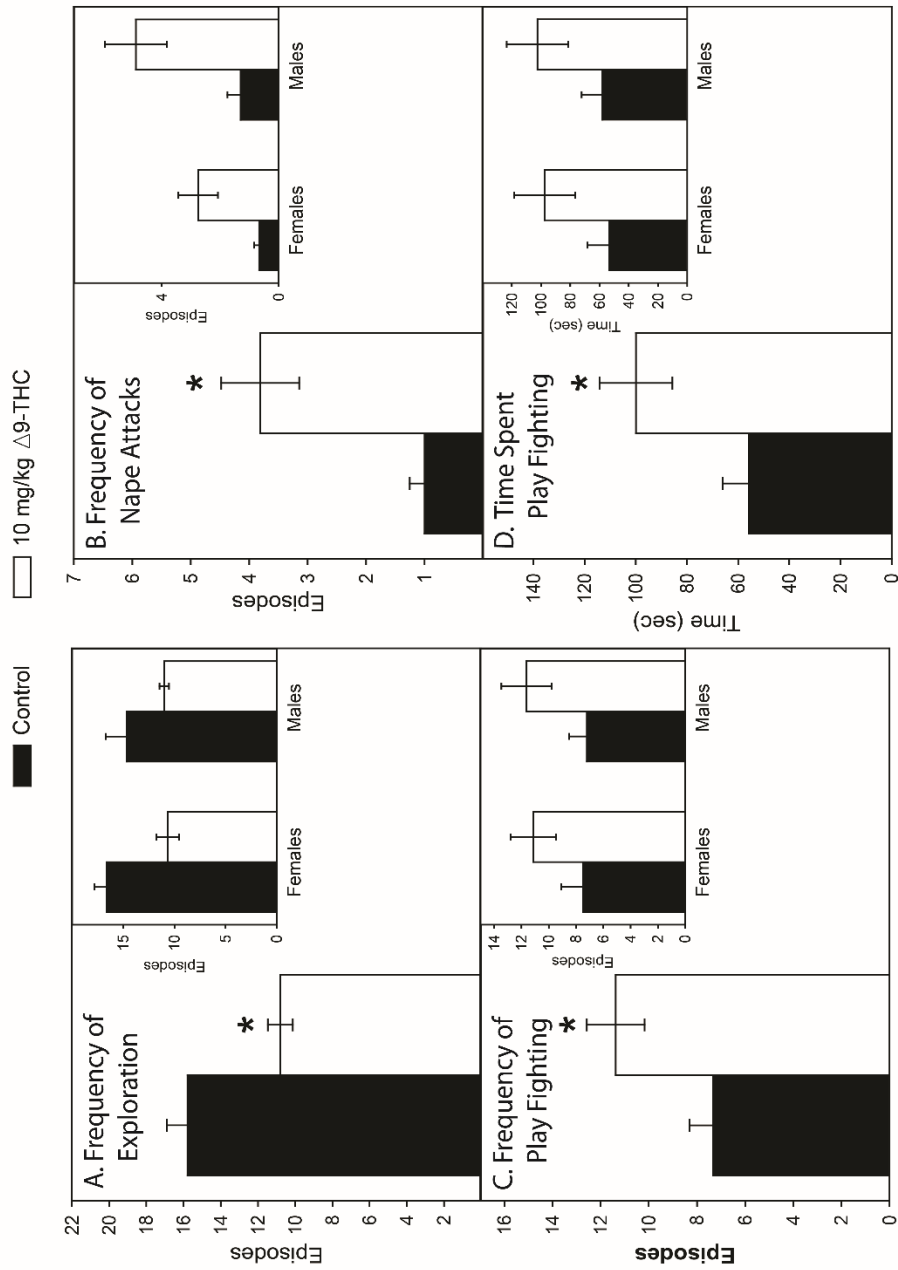


Figure 2.4 Social behavior interactions as described in the Materials and Methods. Frequency of social exploration (A), frequency of nape attacks (B), frequency of play fighting (C), and time spent play fighting (D) in adolescent rats (PND38) exposed to either corn oil or 10 mg/kg of  $\Delta^9$ -THC as juveniles from postnatal days 10-16. Insets are values separated by sex. Values are expressed as means  $\pm$  SEM (n=10-12). Bars with asterisks (\*) are statistically significant ( $p \leq 0.05$ ) from controls.

## 2.4 Discussion

The results of this study suggest that  $\Delta 9$ -THC administered during the postnatal period altered anxiety and social behaviors in adolescent rats. The first interesting observation was that  $\Delta 9$ -THC decreased the expected weight gain in postnatal rats. Importantly, the decreased weight gain was not accompanied by any signs of toxicity. Previous studies have also observed decreased weight gain in rat pups exposed postnatally to  $\Delta 9$ -THC (O'Shea et al., 2004). It has been demonstrated that short term exposure of  $\Delta 9$ -THC to naïve humans and animals induces increased sleep (Babson et al., 2017; Angarita et al., 2016). It is possible that the short term exposure to  $\Delta 9$ -THC induced increased sleep resulting in decreased feeding behavior. Another hypothesis may be the dose and route of exposure used in present study may have significantly contributed to reduced weight. The fact that  $\Delta 9$ -THC being an appetite stimulant as reported by other studies holds true only at lower doses. Higher doses on the other hand are responsible for decrease in the food intake and body weight (Cota et al., 2003).

The elevated plus maze (EPM) is a commonly used model to measure the level of anxiety in rats and mice and the illumination of the maze plays an important factor. Combined increases in the percentage of open arm entries and percentage of time spent in the open arms entries are indications of anxiolytic behavior (Bradley et al., 2007; Pellow et al., 1985), and increased ethological risk assessment behaviors, such as head dipping and stretch attended postures, have been proposed to be sensitive measures related to anxiety in the elevated plus maze (Cruz et al., 1994; Rodgers et al., 1999). However, we did not

observe effects on any parameter that would suggest increased or decreased anxiety levels in adolescent rats exposed to  $\Delta$ 9-THC as juveniles. While not directly comparable, O'Shea et al. (2006) reported no effect on anxiety in pubertal rats following early postnatal exposure to the cannabinoid receptor agonist CP 55,940. In other types of developmental exposures, increased anxiety has been reported in adults exposed either during the gestational/lactational period or the pubertal period to  $\Delta$ 9-THC (Keeley et al., 2015; Campolongo et al., 2008) while decreased anxiety has been reported in adults exposed as adolescents (Rubino et al., 2008; Wakeford et al., 2016). Thus, a direct link between developmental activation of the endocannabinoid system and long term effects on anxiety levels is not straightforward.

In the EPM, multiple studies have reported that increasing the level of illumination decreases activity in the open arms (Garcia et al., 2005; Pereira et al., 2005). Although not directly compared, the pattern of our data are in agreement with this line of thought in that regardless of treatment, the rats tested under low illumination had higher numbers of entries into the open arms and spent more time in the open arms than did the rats tested under high illumination. It has also been suggested that anxiogenic effects of drug exposures are more likely to be detected under low-intensity lighting, whereas anxiolytic effects of drug exposures are more likely to be detected under higher intensity lighting (200-400 Lux or more; Leo & Pamplona 2014). Exposure of juvenile rats to  $\Delta$ 9-THC appears to induce sex specific effects in adolescents. The sex specific differences in open arm activity were observed under high illumination, but totally disappeared when the testing was performed under low illumination suggesting that the effects of treatment only emerge during aversive conditions. Under these conditions, control males did

exhibit a slightly higher open arm activity than did control females but not enough to be statistically significant. Treatment with  $\Delta 9$ -THC did not alter female behavior but exacerbated the open arm activity in males such that sex differences were detected between  $\Delta 9$ -THC males and females. The basis for these sex differences could be due to the differential effect of  $\Delta 9$ -THC exposure on CB1 receptor function between in males and females. Differences exist in CB1 receptor density between males and females in the early postnatal period (Rodriguez de Fonseca et al., 1993). Adolescent exposure to  $\Delta 9$ -THC has been reported to induce sex-differences in the intensity of CB1 receptor-G protein interactions and increased altered CREB phosphorylation (Rubino et al., 2008). It is possible that  $\Delta 9$ -THC exposure during the postnatal period could induce sex differences in CB1 signaling levels as well.

An increase in the number of closed arm entries is an index of locomotor activity (Hogg 1996) and regulating locomotor activity is one of the physiological functions of CB1 receptors in adolescents (Fride 2002). Increased closed arm entries were observed in  $\Delta 9$ -THC treated males, but not  $\Delta 9$ -THC treated females, regardless of the level of illumination. We also observed increased head dipping under both high (increased but not statistically significant) and low illumination. While increased head dipping has been associated with anxiety levels, the increase observed in this study is more than likely merely an artifact of the overall level of increased activity. Increased spontaneous behavior and decreased habituation has also been reported in adult mice exposed to 10 mg/kg  $\Delta 9$ -THC as neonates (Philoppot et al., 2016). However, other studies have reported no effects on activity in adults exposed to  $\Delta 9$ -THC either prenatally or as

adolescents (Newsome & Kelly 2008; Rubino et al., 2008) indicating that, the exposure period and age of testing plays a role in the long term behavioral effects.

Social play is the earliest form of non-mother directed behavioral contact and is crucially important for the normal development of social and cognitive function (Vanderschuren et al., 1997). This behavior is observed to the greatest extent in adolescent rats as compared to juveniles and adults and represents the characteristic social interaction between adolescent rats (Trezza & Vanderschuren 2008). While the actual neural mechanisms involved in social play remain to be elucidated, it has been demonstrated that social play is strongly influenced by the endocannabinoid system in the brain and can be modulated by drugs that directly or indirectly affect activation of the CB1 receptors (Trezza et al., 2012; Trezza & Vanderschuren 2008). In fact, developmental exposure to CB1 receptor agonists can have long term effects on social behavior but the results vary based on the age of treatment, the duration of treatment, and the age of testing. For example, gestational/lactational  $\Delta$ 9-THC exposure did not alter social exploration but decreased social play in adolescents (Trezza et al., 2008) while gestational  $\Delta$ 9-THC exposure increased total social interactions (all social parameters combined) in adults (Newsome & Kelly 2008). Adolescent exposure to  $\Delta$ 9-THC has been reported to either decrease total social investigation (Realini et al., 2011; Zamberletti et al., 2012) or have no effect (Quinn et al., 2008) in adults. In the present study, juvenile exposure to  $\Delta$ 9-THC resulted in a lower level of social investigation in adolescent rats but an increase in social play. At face value, it seems unlikely that rats that exhibit lower levels of investigative interactions would actually have increased levels of playful interactions. One possible explanation could be that since the  $\Delta$ 9-THC treated rats exhibited a higher propensity for

pouncing/nape attack which led to a higher propensity for rough and tumble play, the level of social contact required prior to initiation of play behavior was reduced as compared to controls. However, many consider social exploration behavior to be not related to social play behavior and have suggested that these two parameters differ with respect to both their ontogeny and their regulation by neurological systems (Trezza et al., 2010; Vanderschuren et al., 1997; Spears 1999). The decreased social exploration could be the result of the fact that the  $\Delta$ 9-THC treated rats spent more time engaged in play fighting which reduced the opportunities for social exploration to occur.

Since social play can become very rough and tumble, it could be construed as a sign of adult-like aggressive behavior especially during the late stages of adolescence. However, it has been suggested that the rough and tumble play in developing rats is controlled by a separate neural system than the aggressive interactions observed in adult rats (Panksepp & Burgdorf 2003; Peills & Peills 2007). While there are conflicting reports about the  $\Delta$ 9-THC exposure leads to aggression, it does have a profound effect on the brain reward circuit (Katsidoni et al., 2013; Lupica et al., 2004). Social play induces a rewarding effect and is modulated by the dopaminergic, noradrenergic, opioid, and endocannabinoid systems, all of which have been implicated in reward (Achterberg et al., 2016; Manduca et al., 2015; Trezza et al., 2010; Trezza & Vanderschuren 2008; Vanderschuren et al., 2016). Thus, the modulation in social play observed in this study may involve discoordination between these different signaling systems. The disruption of the function of a single system during development could affect the appropriate functioning of the other systems at the time of behavioral testing. Regardless of the system responsible for



the altered behavior, this observed dysfunction is the result of the action of  $\Delta 9$ -THC on its primary target, the CB1 receptors, during the exposure period.

The CB1 receptors are responsible for maintaining the excitatory inhibitory balance in the brain by controlling the release of glutamate and  $\gamma$ -aminobutyric acid (GABA) at their respective synapses (Castillo et al., 2012; Kano et al., 2009). This balance is crucial for regulating and maintaining proper synaptic activity at most of the neurotransmitter systems involved in mediating social behavior. Of interest, van Kerkhof et al. (2013) reported both the activation of the GABAergic signaling in the nucleus accumbens core or the inhibition of the glutamatergic signaling in the dorsomedial striatum increased social play duration similar to that observed in this study. Therefore, a possible hypothesis is that exposure to  $\Delta 9$ -THC altered the appropriate developmental events necessary for maintaining the excitation-inhibition balance between glutamatergic and GABAergic signaling in specific regions important in social behavior. Development of the GABAergic inhibitory synapses occurs postnatally prior to adolescence in rodents (De Felipe et al., 1997) suggesting that this development might be a target.

In conclusion, given the packaging used in many cannabis edibles (MacCoun et al., 2015), it is not surprising that children find them very enticing. However, the dosages found in these products are highly variable and unregulated but the psychotropic effects are significantly more persistent than other routes of intake (Grotenhermen 2003) suggesting prolonged activation of the neurological target. This is of significant concern with respect to developmental exposures. This research provides an initial investigation into childhood oral exposure to  $\Delta 9$ -THC, an exposure scenario that has recently dramatically increased in occurrence. The results presented here suggest that repeated

oral exposure to  $\Delta$ 9-THC during postnatal development has no effect on anxiety, but can potentially affect the locomotor activity, and social behaviors during adolescence.

However, it is not clear if these are accompanied by increased levels of other behaviors (i.e., risk-taking and novelty-seeking) that are characteristic of adolescent ages (Spears 1999). It is also not clear if exposure to lower dosages will also exert similar effects. This study was limited to a single dosage due to the amount of  $\Delta$ 9-THC available for study.

Additional studies are required to investigate these issues as well as determine the mechanisms responsible for the altered behavior observed in the present study.

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CHAPTER III  
ALTERATIONS IN THE PROTEOMIC PROFILE AND CANONICAL PATHWAYS  
IN ADOLESCENT RATS EXPOSED TO  $\Delta$ 9-TETRAHYDROCANNABINOL AS  
JUVENILES.

### 3.1 Introduction

With the legalization of marijuana and related products in several states of USA, an increase in the marijuana intoxication in children has been reported (Onders et al., 2016; Wang et al., 2014). The psychoactive component of marijuana is  $\Delta$ 9-tetrahydrocannabinol ( $\Delta$ 9-THC ; Gaoni and Mechoulam 1964) which targets the type I cannabinoid receptors (CB1) in brain. The binding of  $\Delta$ 9-THC to CB1 receptors is responsible for producing various behavioral effects associated with  $\Delta$ 9-THC exposure. CB1 receptors are involved in a variety of functions in adult brains which are different from the functions of CB1 receptors in the developing brains of adolescents and juveniles. This fact is supported by the atypical distribution of CB1 receptors in different brain areas (Berrendero et al., 1998). During the gestational (Berrendero et al., 1998) or postnatal period (Romero et al., 1997), CB1 receptors are transiently detected in brain regions that are not normally associated with CB1 signaling in the adult brain. In addition, CB1 mRNA and protein decrease gradually and progressively in juvenile and adolescent brains to finally reach adult levels (Heng et al., 2011). Considering that juvenile and adolescence are two important stages in terms of brain development,

exposure to  $\Delta$ 9-THC during any of these periods could have a long-lasting impact on the functioning of the adult brain and may increase the likelihood of developing psychiatric disorders (Renard et al., 2014).

Previously, adult studies have addressed the effect of  $\Delta$ 9-THC exposure on the expression of different parameters such as tyrosine hydroxylase (Bosier et al., 2012), dopamine receptors (Ginovart et al., 2012), oxytocin receptors (Butovsky et al., 2006), brain derived neurotropic factor (BDNF; ElBatsh et al., 2012. Butovsky et al., 2005), glutamate receptor subunits (Fan et al., 2010), the transcription factor cFOS (Marchese et al., 2008, McGregor et al., 1998, Miyamoto et al., 1997), orexin receptors (Flores et al., 2016), the synaptic vesicle protein VAMP2, and the postsynaptic protein PSD95 (Rubino et al., 2009). However, these studies reported the effects on these parameters in various brain regions immediately following  $\Delta$ 9-THC exposure. Other studies have addressed the issue of persistent behavioral and biochemical effects following  $\Delta$ 9-THC exposure in adolescents or adults (Kasten et al., 2017, Murphy et al., 2017, Weed et al., 2016, Zamberletti et al., 2016;2015;2014, Fishbein et al., 2012). However, the literature is limited with respect to addressing the adverse outcome of early onset of cannabis use on adolescents. We have previously reported that social behavior is altered in adolescents following repeated juvenile exposure to  $\Delta$ 9-THC (Mohammed et al. 2018). The present study focused on determining the changes induced by this repeated juvenile exposure to  $\Delta$ 9-THC on the proteomic profile in the amygdala of adolescent rats. The amygdala was selected because it is one of the brain regions that regulates social behavior (Trezza et al.,

2013) and because it contains a high density of CB1 receptors (Herkenham et al., 1990), making it one of the brain regions that is a likely target for  $\Delta$ 9-THC-mediated effects (Bindukumar et al., 2008, Renard et al., 2014). In this study, label free proteomic analysis was conducted to determine changes in protein expression followed by further analysis to determine the pathways and functions most strongly affected by juvenile  $\Delta$ 9-THC exposure.

## **3.2 Materials and Method**

### **3.2.1 Chemicals**

$\Delta$ 9-tetrahydrocannabinol was provided by the National Institute on Drug Abuse. All other chemicals were purchased from Sigma Chemical Co. (St. Louis, MO).

### **3.2.2 Animals**

A breeding colony of adult male and female Sprague Dawley rats (CD IGS; Harlan Laboratories, Indianapolis, IN) were housed in an Association for Assessment and Accreditation of Laboratory Animal Care-accredited facility with temperature control ( $22 \pm 2^\circ\text{C}$ ) and with lights on between 0700 and 1900 for a 12-h dark-light cycle. Tap water and LabDiet rodent chow were freely available during the experimentation. The procedures used in this project were approved by the Mississippi State University Institutional Animal Care and Use Committee. From this colony, male and female rat pups were obtained and the day of birth was considered as PND0. On PND10, rat pups within the same litter were assigned to different treatment groups. There was always a representative control animal of the same sex present in each litter to match the  $\Delta$ 9-THC-treated animal.

### **3.2.3 Exposure**

Beginning on PND10, rats were exposed once a day for 7 days to either corn oil (vehicle) or 10 mg/kg  $\Delta^9$ -THC by oral gavage at a volume of 0.5 ml/kg. Solutions were delivered to the back of the throat using a 50- $\mu$ l tuberculin syringe equipped with a 1-inch 24-gauge straight intubation needle (Popper and Sons, Inc., New Hyde Park, NY). Rats were weaned on PND21 and remained in cages undisturbed until PND38.

### **3.2.4 Proteomic analysis**

On the PND38, the whole brain was collected, frozen on a stainless steel plate on top of dry ice, and stored at  $-80^{\circ}$  C. Three biological replicates of control and  $\Delta^9$ -THC-treated male rats from different litters were used. Each brain was subjected to coronal sectioning to obtain 500-micron thick brain slices. Using the rat brain atlas as a reference (Pixons and Watson 1998), the amygdala was isolated using 1 mm punch and lysed in NP-40 lysis buffer. The protein concentration was measured using a Pierce™ BCA protein assay kit (Thermo Scientific). From each sample, 100  $\mu$ g of protein was precipitated by chloroform/methanol extraction. After precipitation, protein digestion was performed by suspending the pellet in 33  $\mu$ l of 100 mM Tris-HCl (pH 7.8) containing 6 M urea. The samples were reduced with 1.6  $\mu$ l of 200 mM dithiothreitol (DTT) followed by alkylation with 6.6  $\mu$ l of 200 mM iodoacetamide (IAA) each for 45 min at room temperature. The alkylation reaction was quenched by adding 20  $\mu$ l of 200 mM DTT for 45 min at room temperature. The urea concentration was reduced by adding 258  $\mu$ l of milliQ-H<sub>2</sub>O. Finally, the proteins were digested with trypsin at 1:50 ratio for 18 hr at 37°C. Protein digestion was terminated by lowering the pH of each sample to <6 by adding concentrated acetic acid. The samples were desalted using C18 SepPak columns (Waters,

USA). These samples were submitted to the University of Arizona Proteomics Consortium for analysis of samples by using Tandem mass spectrometry coupled to liquid chromatography (LC-MS/MS).

The LC-MS/MS analysis of trypsin digested protein samples was performed by using LTQ Orbitrap Velos mass spectrometer (Thermo Fisher Scientific, San Jose, CA) equipped with an Advion nanomate ESI source (Advion, Ithaca, NY). Peptides were eluted from a C18 pre-column (100- $\mu$ m id  $\times$  2 cm, Thermo Fisher Scientific) onto an analytical column (75- $\mu$ m ID  $\times$  10 cm, C18, Thermo Fisher Scientific) using a beginning concentration of 2% solvent B (acetonitrile, 0.1% formic acid) for 5 minutes, then a 2–7% gradient of solvent B over 5 minutes, followed by a 7-15 % gradient of solvent B over 50 minutes, a 15-35% gradient of solvent B over 60 minutes, a 35-40% gradient of solvent B over 28 minutes, a 40-85% gradient of solvent B over 5 minutes, held at 85% solvent B for 10 minutes, 85-2% gradient of solvent B for 1 minute then held at 2% solvent B for 16 min. All flow rates were at 400 nl/min. Solvent A consisted of water and 0.1% formic acid. Data dependent scanning was performed by the Xcalibur v 2.1.0 software (Andon et al., 2002) using a survey mass scan at 60,000 resolution in the Orbitrap analyzer scanning m/z 400–1600, followed by collision-induced dissociation (CID) tandem mass spectrometry (MS/MS) of the fourteen most intense ions in the linear ion trap analyzer. Precursor ions were selected by the monoisotopic precursor selection (MIPS) setting with selection or rejection of ions held to a  $\pm$  10 ppm window. Dynamic exclusion was set to place any selected m/z on an exclusion list for 45 seconds after a single MS/MS. All MS/MS spectra were searched against a protein database using



Thermo Proteome Discoverer 1.3 (Thermo Fisher Scientific, San Jose, CA) using a Rattus Norvegicus protein database downloaded from Uniprot (<http://www.uniprot.org>). Fully tryptic peptides with up to 2 missed cleavage sites were selected. Variable modifications considered during the search included methionine oxidation (15.995 Da), and cysteine carbamidomethylation (57.021 Da). Proteins were identified at 99% confidence with XCorr score cut-offs (Qian et al., 2005) as determined by a reversed database search.

### **3.2.5 Data analysis**

The proteins and peptide identification results were visualized by using scaffold v 3.6.1 (Proteome software Inc., Portland, OR, USA). Protein identification with a minimum of two peptides identified with a 0.1% false discovery rate (FDR) were deemed correct. Significant changes in the protein expression was calculated by using Fisher's exact test. Fold changes in the protein expression were calculated using weighted normalized spectra with 0.2 imputation value.

### **3.2.6 IPA and gene ontology**

For pathway and ontology analysis, the protein list was filtered by applying the LOG2FC threshold of 0.6 and the Fisher's exact P-value of 0.08 to ensure the presence of closely related and biologically significant proteins in the list. Biological interpretation of identified proteins was performed by DAVID (Database of Annotation, Visualization and Integrated Discovery). The functional annotation clustering categorizes the differentially expressed proteins among the controlled vocabulary of Gene Ontology (GO) in three main categories: Biological process, which describes broad biological goal;

Molecular function, which describes the task performed by individual gene products; and Cellular component, which describes the location of molecules inside the cell (Dennis et al., 2003). In current study, GO terms enriched with differentially expressed protein with a P-value  $\leq 0.05$  were selected.

Differentially expressed proteins were also analyzed in Ingenuity Pathway Analysis (IPA; Qiagen Inc., <https://www.qiagenbioinformatics.com>) to obtain the canonical pathways, physiological functions, and networks associated with these proteins. In the initial analysis for pathways in IPA, only differentially expressed peptides were included that passed the LOG2FC cutoff 0.6, and Fisher's exact P-value of 0.05 were included which identified some biologically relevant pathways. A subsequent analysis was performed using a Fisher's exact P-value of 0.08 to include more differentially expressed proteins. Addition of proteins did not change the enrichment of pathways but improved the z-scores of pathways that were enriched at P-value of 0.05 (Figure 1).

### **3.2.7 Quantitative PCR (QPCR)**

Total RNA from the amygdala was extracted using the Qiagen RNeasy mini kit (Germantown, MD), and converted to cDNA by using the Qiagen RT2 cDNA synthesis kit provided with genomic DNA elimination buffer (Germantown, MD). Pre-designed gene specific primers, probes (Table 3.1), and Primetime Gene Expression master mix (Integrated DNA Technology, Coraville, IA) were used to perform real time QPCR. Each sample was run in triplicate in a final reaction volume of 20 ul according to manufacturer's instructions. The thermal cycler conditions were as follows: polymerase activation- 95° C for 3 min, amplification- 40 cycles of 95° C for 15 sec and 60° C for 1 min. In each experiment, standard curves for the gene of interest and the reference gene

were made by using a range of cDNA dilutions (1:5-1:1000) and the PCR efficiency factor was calculated. The threshold Ct value of the gene of interest were normalized with Ct values of the reference gene and PCR efficiency factor. Normalized Ct values were used to calculate the fold change from control by using method  $2^{(Ct(\text{control mean}) - Ct(\text{Treatment}))}$  as described by Betancourt et al. (2007). Statistical analysis was conducted on fold change values by using non-parametric Kruskal-Wallis one-way analysis of variance to determine the significant treatment effect. The level of statistical significance was set at  $P \leq 0.05$ .

Table 3.1 Primers, probes, and assay names for reference and target genes

Gene name	Assay name	Sequence 5'-3'
Actb	Rn.PT.	Primer 1 GGCATAGAGGTCTTTACGGATG
	39a.222214	Primer 2 TCACTATCGGCAATGAGCG
	838.g	Probe  /5HEX/TCCTGGGTA/ZEN/TGGAATCCTGTGGC/3IAkFQ
Gnai	Rn.PT.58.5	Primer 1 GTCGTTTTCACTCTAGTTCTGAGA
	104082	Primer 2 GAGTACCAGCTGAACGATTCG
		Probe  /56FAM/CAACCAAAT/ZEN/TACATCCCAACCCAGCA3IABk  FQ

Table 3.1 (continued)

Gnas	Rn.PT.58.6 088247	Primer 1 CCTTCTCACCATCGCTGTT Primer 2 CACCATTGTGAAGCAGATGAG Probe /56FAM/CCGCCCTCT/ZEN/CCGTAAACCCATTAA/3IABkF Q
Gnaq	Rn.PT.58.3 8203772	Primer 1 CACAAGCTCTTTATTGCGTCTA Primer 2 CACTCTCAAGATCCCATAACAAGT Probe 56FAM/ACACCTTCT/ZEN/CCACATCAACCTCTCGA/3IABkF Q
Gna13	Rn.PT.58.3 4054555	Primer 1 GCCATCAACTTGTCTCCGT Primer 2 CACCATCTACAGCAACGTGA Probe 56FAM/CTTCTCTCG/ZEN/GGCATCTACCAGCAC/3IABkF

### 3.3 Results and Discussion

This study was designed to determine the persistent changes in amygdaloid proteomic expression in adolescent rats that occur as a result of repeated exposure to  $\Delta^9$ -THC at

juvenile stage. The differential expression in proteins due to  $\Delta 9$ -THC exposure was determined by applying Fisher's exact P-value threshold ( $P \leq 0.08$ ), and LOG2FC threshold of 0.6. Out of a total of 1169 proteins detected, there were 38 proteins that were up regulated and 52 proteins that were down regulated due to juvenile  $\Delta 9$ -THC exposure. Functional clustering of differentially expressed proteins in DAVID revealed that most of the differentially expressed proteins were part of the cellular components with GO terms such as synapse, presynaptic membrane, synaptic vesicle and post-synaptic density significantly enriched (Figure 2) by differentially expressed proteins.

In juveniles, atypical distribution of the CB1 receptors in different brain regions is responsible for guiding the elongating axons to their targets in order to form a synapse with the postsynaptic neurons in the target region (Harkany et al., 2008). Large numbers of differentially expressed synaptic proteins in our study suggest that the improper CB1 activation during juvenile  $\Delta 9$ -THC exposure affected the axonal guidance cue which disrupted appropriate synapse formation. Altered synapse formation could be the molecular basis for the social play deficits that we observed in our previous study (Mohammed et al., 2018).

Table 3.2 lists the total canonical pathways enriched in IPA analysis of differentially expressed proteins in this study. There are some relevant pathways with positive or negative activity score are opioid signaling, G $\alpha$ q signaling, Rho family GTPase, RhoGDI signaling, and thrombin signaling (Figure 3). The differentially expressed proteins associated with these pathways are listed in Table 3.3.

Table 3.2 Canonical pathways predicted using differentially expressed proteins in the amygdala of adolescent rats (PND38) exposed to either corn oil or 10 mg/kg  $\Delta$ 9-THC during juvenile period (PND10-16).

Canonical pathways	Activity z-score
RhoGDI signaling	Negative
Signaling by Rho family GTPase	Positive
Protein ubiquitination pathway	No pattern
Thrombin signaling	Positive
CXCR4 signaling	Zero
CREB signaling in neurons	Zero
Ephrin signaling	Zero
Ethanol degradation II	No pattern
Noradrenaline and adrenalin degradation	No pattern
Gaq pathway	Zero
Tec kinase signaling	Zero
G beta-gamma signaling	Zero
IL-B signaling	Zero
Semaphorin signaling in neuron	No pattern
Breast cancer regulation by stathmin	No pattern

Table 3.2 (continued)

Actin nucleation by ASP-WASP complex	Zero
Serotonin degradation	No pattern
CCR3 signaling in eosinophile	Zero
Androgen signaling	Zero
Histamine degradation	No pattern
Cardiac hypertrophy signaling	Zero
Oxidative ethanol degradation III	No pattern
Chemokine signaling	Zero
Huntington disease signaling	No pattern
Fatty acid oxidation	No pattern
GM-CSF signaling	Zero
Putriscine degradation	No pattern
Ethanol degradation IV	No pattern
Choline degradation	No pattern
D-Mannose degradation	No pattern
Tryptophan degradation	No pattern
Alpha-adrenergic signaling	Zero
CCR5 signaling in macrophages	Zero
Ephrin receptor signaling	Zero
IL-1 signaling	Zero

Table 3.2 (continued)

Dopamine degradation	No pattern
G protein signaling mediated by Tubby	No pattern
Cholestokinin/Gestin mediated signaling	Zero
Formaldehyde oxidation II	No pattern
Glutamate dependent acid resistance	No pattern
Oxidative phosphorylation	No pattern
Axonal guidance signaling	No pattern
Neuropathic pain signaling in dorsal horn neurons	Zero
Glioma signaling	Zero
Role of NFAT in cardiac hypertrophy	Zero
Mechanisms of viral exit from host cell	No pattern
Rho A signaling	Zero
EIF2 signaling	Negative
Synaptic long term potentiation	Zero
Phospholipase C	Zero
fMLP signaling	Zero
nNOS signaling in neurons	Zero



Table 3.2 (continued)

Pentose phosphate pathway	No pattern
Opioid signaling	Positive
Phagosome mutation	No pattern
Protein Kinase A signaling	Zero
Lysine degradation II	No pattern
Glutamate degradation III	No pattern
Xenobiotic metabolism	No pattern
Molecular mechanism of cancer	No pattern
Relaxin signaling	Zero
GDP mannose biosynthesis	No pattern
Synaptic long term depression	Zero
Mitochondrial dysfunction	No pattern
GDP-glucose biosynthesis	No pattern
Melatonin signaling	Zero
Role of NFAT regulation in Immune response	Zero
Glucose and glucose 1 phosphate degradation	No pattern
Antiproliferative role of somatostatin receptor 2	Zero
Pentose phosphate pathway	No pattern

Table 3.2 (continued)

UDP-N-acetyl D-galactosamine biosynthesis	No pattern
Purine nucleotide de novo biosynthesis	No pattern
Glycogen degradation	No pattern

### 3.3.2 Gaq signaling

The CB1 receptors are G-protein coupled receptors and their functioning receptors occurs by binding to the Gai subunit leading to the inhibition of adenylyl cyclase (See review: Howlett 2005). This inhibitory action induced by the CB1 receptors is reversed in the presence of pertussis toxin which blocks the binding of the Gai subunit to the CB1 receptor allowing the synthesis and accumulation of cAMP (Felder et al., 1998). Several studies have reported that a stimulatory action can be induced by the CB1 receptors and this occurs through the binding of alternate subunits such as Gas or Gaq depending upon the relative abundance of each subunit (Bonhaus et al., 1996; Glass and Felder 1997; Felder et al., 1998; Lauckner et al., 2005). In the current study, we observed an enrichment of the Gaq signaling pathway which could indicate that the repeated exposure to  $\Delta 9$ -THC persistently altered the abundance of Gai relative to other subunits. This hypothesis was tested by measuring the level mRNA expression of the Gai, Gas, Gaq, and Ga13 subunits using QPCR. The results indicated that there was a 15% fold increase in the mRNA levels of Gai and Gaq whereas the levels of Gas and Ga13 were unaffected (Figure 4). Previously, it has been reported that down regulation of the mRNA expression

for the G protein subunits occurs immediately following repeated exposure to a CB1 agonist (Rubino et al., 1997). Our gene expression data suggest that a reversal of the effect on the gene expression of G protein subunits occurs after cessation of  $\Delta 9$ -THC exposure. However, this data may not explain the enrichment of G $\alpha$ q signaling as seen in our IPA data because the binding of subunit to receptor may not be correlated to the amount of mRNA present in tissue (Rubino et al., 1997).

### **3.3.3 Opioid signaling**

In previous studies, it was reported that behavioral and pharmacological effects induced by  $\Delta 9$ -THC involves co-activation of cannabinoid and opioid receptors (Manharaz et al., 1999; Ambrosio et al., 1999). This synergistic action of cannabinoid and opioid receptors is majorly seen with the reward behaviors and antinociception. Absence of either CB1 (Martin et al., 2000) or opioid receptor (Ghozland et al., 2002) abolished the reward behavior induced by the respective receptor agonist. Chronic exposure to  $\Delta 9$ -THC can enhance the antinociceptive potency of morphine which is an opioid receptor agonist (Cichewicz et al., 1999; Welch and Eads, 1999; Smith et al., 1998).  $\Delta 9$ -THC is also known to produce analgesic effects and one of the major hypotheses behind  $\Delta 9$ -THC-induced analgesia is that it involves synergistic actions of both cannabinoid and opioid signaling (Welch and Eads, 1999). In addition, it has been reported that CB1 and opioid receptors are co-localized in amygdala (Novarro et al., 2001) where activation of CB1 receptor is responsible for producing  $\Delta 9$ -THC-induced analgesia (Lee et al., 2011). Thus, the activation of the CB1 receptor by  $\Delta 9$ -THC can in turn activate opioid signaling. Based on our data, it appears that this activation of the CB1 receptors during juvenile brain development leads to a persistent enhancement of opioid signaling which lasts

through adolescence. One hypothesis behind this  $\Delta 9$ -THC-induced long term alteration in opioid signaling may be that repeated  $\Delta 9$ -THC exposure in juvenile rats has increased the opioid receptor density in amygdala as seen in previous study (Cochero et al., 2004). This may lead to persistent alterations in opioid signaling as seen in our data.

### **3.3.4 Rho GTPase and thrombin signaling**

The Rho family GTPase belong to the Ras superfamily of small peptides. Typically, Rho peptides are cytoskeleton proteins and involved in the organization of microtubules (Gundersen et al., 2004) and formation of stress fibers at the cell terminal (Nobes and Hall 1995) which in turn governs the cell migration and differentiation. In terms of neuronal development, the members of the Rho family are associated with various aspects of neural development such as axon formation, neuronal migration, and neurite outgrowth (Watabe-Uchida et al., 2006). Neurite formation and inhibition are two important phenomena during brain development in order to achieve proper connectivity between different neurons (Skaper 2005). Several studies reported that activation of Rho GTPase negatively regulates in the axonal outgrowth (Gu et al., 2013; Harkany et al., 2008; Ng and Luo 2004; Lehman et al., 1999). Thrombin signaling is also associated axonal growth. Thrombin signaling occurs through G protein coupled receptors and is directly related to neurite retraction both during injury and during normal brain functioning. The addition of *Clostridium botulinum C3 exoenzyme*, which inhibits Rho, blocks the thrombin-induced neurite retraction (Jalink et al., 1994) indicating collaborative action of thrombin signaling and Rho GTPase to control the axonal growth. As Rho GTPases cycle from the cytoplasmic inactive form to the membrane bound form where they are active to interact with their targets. RhoGDP-dissociation inhibitors

(RhoGDI) are regulatory peptides that control Rho signaling by binding to the Rho proteins in the inactive form (Rho-GDP complex) in cytoplasm thus restricting the translocation to the membrane where they are activated (Rho-GTP; Robbe et al., 2003). In our data, there was an inhibition of RhoGDI signaling and activation of Rho GTPase, and thrombin. Thus, these data are suggesting that the juvenile exposure to  $\Delta 9$ -THC has persistent effects on neuronal functioning such as axonal formation, and neuronal migration which are governed by Rho GTPase. Since the juvenile period is a period where synaptogenesis peaks (Casey et al., 2000) and CB1 receptors were detected in the elongating axons that would reach the target to form a synapse (Fernandez-Ruiz et al., 2000),  $\Delta 9$ -THC -mediated activation of CB1 receptor during juvenile period may alter the synaptogenesis by activating Rho GTPase and thrombin signaling.

### **3.4 Conclusion**

It is well established that the endocannabinoids (EC) and the CB1 receptors are involved in antinociception and neuronal development. Activation of opioid signaling in our data may validates a previous hypothesis that antinociceptive action of THC is mediated through amygdala. Initially, studies only predicted their functions in neural development based on the presence of CB1 receptors on developing neurons in specific regions (Romero et al., 1997). Later, involvement of the CB1 receptors in synaptogenesis was proposed (Fernandez-Ruiz et al., 2000). Our data has provided insight about how the  $\Delta 9$ -THC exposure during postnatal ages could persistently affect the neuronal pathways that are negatively correlated with in axonal migration, or neuritogenesis. As these processes are eventually involved in synaptogenesis, it is possible that juvenile  $\Delta 9$ -THC exposure may affect the synaptogenesis in postnatal brain. This is also supported by the functional



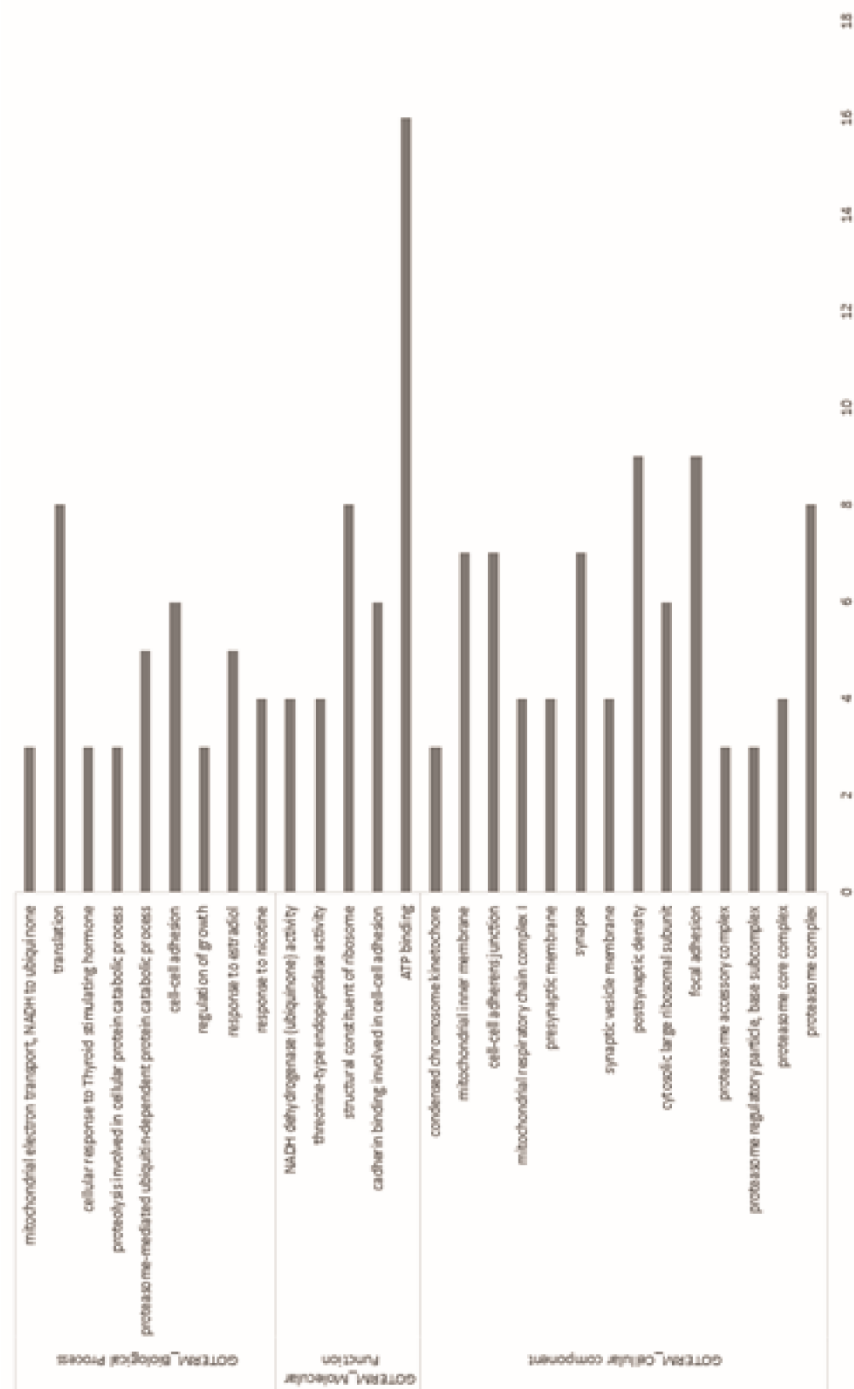


Figure 3.2 GO term analysis of differentially expressed proteins obtained from amygdala of adolescent rats (PND38) repeatedly exposed to either corn oil or 10 mg/kg  $\Delta^9$ -THC during juvenile ages (PND10-16).

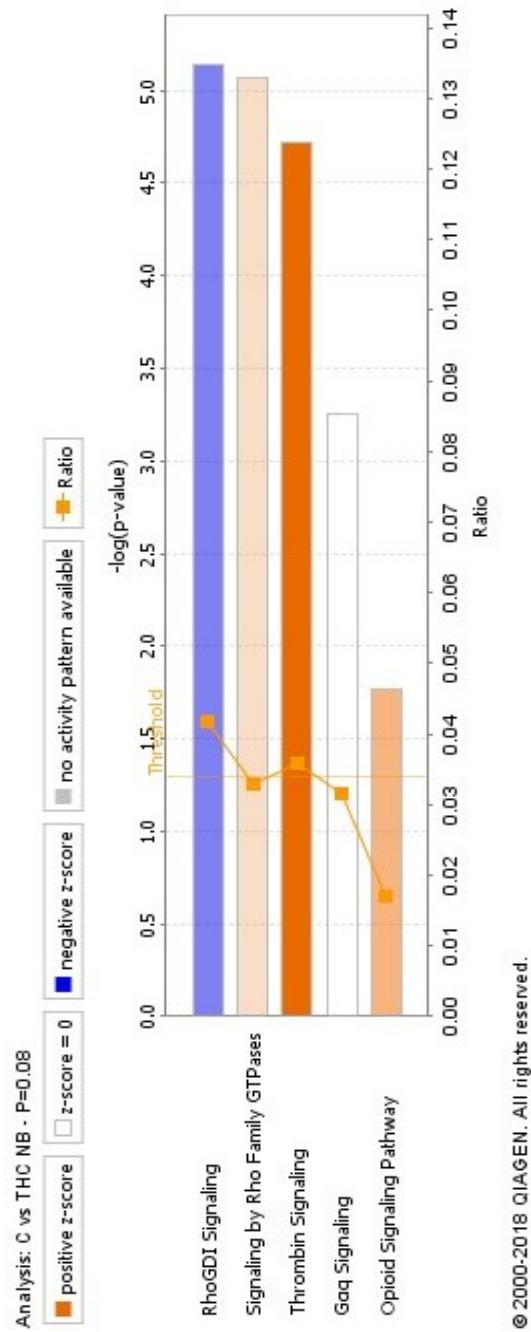


Figure 3.3 Canonical pathways associated with differentially expressed proteins obtained from amygdala (PND38) of rats exposed repeatedly to either corn oil or 10 mg/kg  $\Delta^9$ -THC at juvenile ages (PND10-16).



Table 3.3 List of differentially expressed protein involved in the relevant canonical pathways shown in figure 3.3

Uniprot Accession	Gene Symbol	Entrez Gene name	Regulation
KCC2A_RAT	<i>CAMK2A</i>	Calcium/calmodulin dependent protein	Up
F1M7X3_RAT	<i>CDH3</i>	Cadherine 13	Down
A0A0G2K526_RAT	<i>GNAL</i>	G protein subunit alpha L	Up
G3V8K2_RAT	<i>GNG3</i>	G protein subunit gamma 3	Down
RACK1_RAT	<i>RACK1</i>	Receptor for activated C kinase 1	Down
A0A0G2K5N6_RAT	<i>ROCK2</i>	Rho associated coiled coil containing protein	Up
RHOB_RAT	<i>RHOB</i>	Ras containing family member B	Up
WASF1_RAT	<i>WASF1</i>	Was containing family member 1	Up

Table 3.3 (continued)

FILMH0_RAT	<i>SEPT 3</i>	Septin 3	Down
A0A0G2K526_RAT	<i>PRKB</i>	Protein kinase C Beta	Up
PENK_RAT	<i>PENK</i>	Proenkaphalin	Down

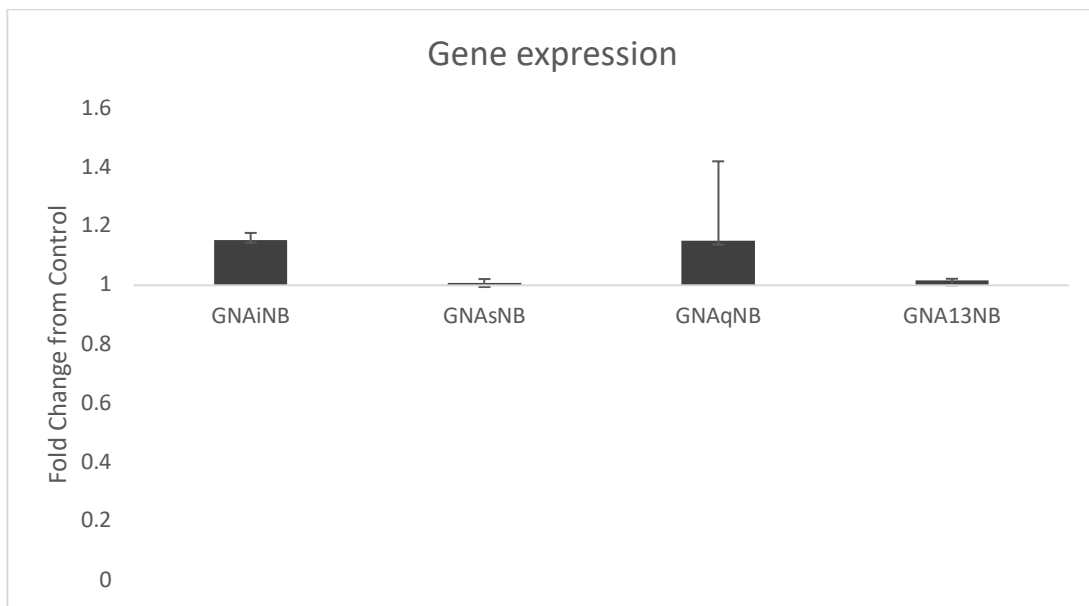


Figure 3.4 Change in mRNA expression in amygdala of adolescent rats (PND38) repeatedly exposed to either corn oil or 10 mg/kg of  $\Delta$ 9-THC during juvenile ages (PND10-16).

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CHAPTER IV  
ALTERATIONS IN CB1-RELATED CANONICAL PATHWAYS DURING SOCIAL  
PLAY IN ADOLESCENT RATS REPEATEDLY EXPOSED TO  $\Delta$ 9-  
TETRAHYDROCANNABINOL AS JUVENILES

**4.1 Introduction**

Social play is one of the earliest form of interactions with peers that occurs without any involvement of the mother. Because of this feature, adolescent social play is included among several important phenomena that are essential for developing normal adult social behaviors (Pellis and Pellis 2009; van den Berg et al., 1998). The occurrence of social play is one of the useful biomarkers for neurodevelopment (Blake and McCoy 2015) and deficits in play performance have been connected to certain neuropsychiatric disorders (Trezza et al., 2010; Meany and Stewart 1981). Play is a characteristic behavior that starts during the early stage of life, peaks during the adolescence, and declines at the time of puberty (Trezza et al., 2010; Martin and Caro et al., 1985). Despite its strong influence in terms of brain development, our knowledge towards understanding the physiological mechanism and neural substrates underlying social play is limited (Blake and McCoy 2015; Trezza and Vanderschuren 2008a). Currently, studies are focusing on elucidating the involvement of the brain endocannabinoid system in mediating social play behavior through type I Cannabinoid receptors (CB1; Trezza and Vanderschuren 2008b).

The modulation of social play through endocannabinoid signaling mainly occurs in the amygdala as demonstrated by Trezza and colleagues (2013). They investigated the effects of activation of endocannabinoid signaling on social play by administering the FAAH inhibitor URB597 that acts as an indirect CB1 agonist and observed that the level of accumulation of endogenous CB1 ligand (anandamide or AEA) was highest in the amygdala as compared to other regions such as the hippocampus and the nucleus accumbens. Another study focused on the role of CB1 ligands AEA and 2-AG (2-arachidonoyl glycerol) in the play behavior. They reported an increase in the social play activity in rats treated with a dual inhibitor of AEA and 2-AG hydrolysis. This increase in the social play activities was antagonized by CB1 receptor antagonist (Manduca et al., 2015).

The CB1 receptors are the primary target of  $\Delta 9$ -tetrahydrocannabinol ( $\Delta 9$ -THC; Devane et al., 1992) which is the main psychoactive component of marijuana (Gaoni and Mechoulam 1964). The CB1 receptors are involved in a variety of functions related to neurogenesis (Berrendo et al., 1998; Romero et al., 1997; Fernandez-Ruiz et al., 2000; Heng et al., 2011) and some of the forms of neurogenesis, such as neuronal migration and synaptogenesis, peak during the juvenile period (Markus and Petit 1987). Thus, juvenile exposure to  $\Delta 9$ -THC could be catastrophic in terms of establishing normal neurological function which mainly depends on synaptogenesis during the early stages of life (Diaz-Alonso et al., 2012). When investigating the  $\Delta 9$ -THC-mediated alterations in brain development, determining the altered canonical pathways in the proteome as it relates to behavioral function could provide very insightful information. Several studies have reported immediate or long-term effects of  $\Delta 9$ -THC exposure on social interactions at

various life stages (Manduce et al., 2015; Trezza et al., 2013; Trezza and Vanderschuren 2008a,b Quinn et al., 2008; Newsome and Kelly 2008; Novarro et al., 1995; Cutler and Mackintosh 1984; van Ree et al., 1984; Seiber 198; Miczek 1974). However, none of these studies have investigated the  $\Delta$ 9-THC-mediated changes in the biochemical pathways related to social play. In our previous study, we reported that adolescent rats receiving  $\Delta$ 9-THC as juveniles engaged in playful activities for longer duration as compared to control adolescent rats (Mohammed et al., 2018). We have also determined the canonical pathways in the proteome altered in adolescents exposed to  $\Delta$ 9-THC as juveniles in the absence of any behavioral activity. In this study, we determined the alterations in the canonical pathways in the proteome related directly to  $\Delta$ 9-THC-mediated changes in social play.

## **4.2 Materials and Method**

### **4.2.1 Chemical**

$\Delta$ 9-tetrahydrocannabinol was provided by the National Institute on Drug Abuse. All other chemicals were purchased from Sigma Chemical Co. (St. Louis, MO).

### **4.2.2 Animals**

A breeding colony of adult male and female Sprague Dawley rats (CD IGS; Harlan Laboratories, Indianapolis, IN) were housed in an Association for Assessment and Accreditation of Laboratory Animal Care-accredited facility with temperature control ( $22 \pm 2^\circ\text{C}$ ) and with lights on between 0700 and 1900 for a 12-h dark-light cycle. Tap water and LabDiet rodent chow were freely available during the experimentation. The procedures used in this project were approved by the Mississippi State University

Institutional Animal Care and Use Committee. From this colony, male and female rat pups were obtained and the day of birth was considered as PND0. On PND10, rat pups within the same litter were assigned to different treatment groups. There was always a representative control animal of the same sex present in each litter to match the  $\Delta$ 9-THC-treated animal.

### **4.2.3 Exposure**

Beginning on PND10, rats were exposed once a day for 7 days to either corn oil (vehicle) or 10 mg/kg  $\Delta$ 9-THC by oral gavage at a volume of 0.5 ml/kg. Solutions were delivered to the back of the throat using a 50- $\mu$ l tuberculin syringe equipped with a 1-inch 24-gauge straight intubation needle (Popper and Sons, Inc., New Hyde Park, NY). Rats were weaned on PND21 and categorized into either a behavioral cohort or a non-behavior cohort. Each cohort consisted of set of rats receiving either vehicle or  $\Delta$ 9-THC.

### **4.2.4 Animal handling**

#### **4.2.4.1 Behavior rats**

After weaning on PND21, rats belonging to the behavioral cohort were subjected to testing in the elevated plus maze (EPM) on PND29 under low illumination followed by testing for social interactions on PND38 (data presented in Chapter 2). On PND37, these rats were individually housed for a 24-hour isolation period. A pair of rats of the same sex and size and belonging to the same treatment group but from different litters were placed into a novel cage under high illumination (600 lux) and allowed to interact for 10-min interval. After social play, the rats were returned to the isolation cage and sacrificed 3-hours later.

#### **4.2.4.2 Non-behavior rats**

Rats from the non-behavior cohort were handled in the same way as behavior rats with the exception that they were not subjected to any type of behavior testing. After weaning on PND21, these rats were individually housed on PND37 for a 24-hour isolation period and then sacrificed 24-hours following isolation.

#### **4.2.5 Proteomic analysis**

On PND38, each rat in both the behavioral and non-behavioral cohorts were sacrificed to collect the whole brain which was frozen on a stainless steel plate on top of dry ice and stored at -80°C. Three biological replicates, following exposure conditions, were used.

- 1) Behavioral Control Rats, which received corn oil and subjected to social play.
- 2) Behavioral  $\Delta$ 9-THC Rats, which received 10 mg/kg of  $\Delta$ 9-THC and subjected to social play.
- 3) Non-behavioral Control Rats, which received corn oil but not subjected to social play.
- 4) Non-behavioral  $\Delta$ 9-THC Rats, which received  $\Delta$ 9-THC but not subjected to social play.

Each brain was subjected to coronal sectioning to obtain 500-micron thick brain slices.

Using the rat brain atlas as a reference (Paxinos and Watson 1998), the amygdala was isolated using 1 mm punch and lysed in NP-40 lysis buffer. The protein concentration was measured using a Pierce™ BCA protein assay kit (Thermo Scientific). From each sample, 100  $\mu$ g of protein was precipitated by chloroform/methanol extraction. After precipitation, protein digestion was performed by suspending the pellet in 33  $\mu$ l of 100 mM Tris-HCl (pH 7.8) containing 6 M urea. The samples were reduced with 1.6  $\mu$ l of 200 mM dithiothreitol (DTT) followed by alkylation with 6.6  $\mu$ l of 200 mM

iodoacetamide (IAA) each for 45 min at room temperature. The alkylation reaction was quenched by adding 20  $\mu$ l of 200 mM DTT for 45 min at room temperature. The urea concentration was reduced by adding 258  $\mu$ l of milliQ-H<sub>2</sub>O. Finally, the proteins were digested with trypsin at 1:50 ratio for 18 hr at 37°C. Protein digestion was terminated by lowering the pH of each sample to <6 by adding concentrated acetic acid. The samples were desalted using C18 SepPak columns (Waters, USA). These samples were submitted to the University of Arizona Proteomics Consortium for Orbitrap LC-MS/MS analysis. The LC-MS/MS analysis of trypsin digested protein samples was performed by using LTQ Orbitrap Velos mass spectrometer (Thermo Fisher Scientific, San Jose, CA) equipped with an Advion nanomate ESI source (Advion, Ithaca, NY). Peptides were eluted from a C18 pre-column (100- $\mu$ m id  $\times$  2 cm, Thermo Fisher Scientific) onto an analytical column (75- $\mu$ m ID  $\times$  10 cm, C18, Thermo Fisher Scientific) using a beginning concentration of 2% solvent B (acetonitrile, 0.1% formic acid) for 5 minutes, then a 2–7% gradient of solvent B over 5 minutes, followed by a 7-15 % gradient of solvent B over 50 minutes, a 15-35% gradient of solvent B over 60 minutes, a 35-40% gradient of solvent B over 28 minutes, a 40-85% gradient of solvent B over 5 minutes, held at 85% solvent B for 10 minutes, 85-2% gradient of solvent B for 1 minute then held at 2% solvent B for 16 min. All flow rates were at 400 nl/min. Solvent A consisted of water and 0.1% formic acid. Data dependent scanning was performed by the Xcalibur v 2.1.0 software (Andon et al., 2002) using a survey mass scan at 60,000 resolution in the Orbitrap analyzer scanning m/z 400–1600, followed by collision-induced dissociation (CID) tandem mass spectrometry (MS/MS) of the fourteen most intense ions in the linear ion trap analyzer. Precursor ions were selected by the monoisotopic precursor selection

(MIPS) setting with selection or rejection of ions held to a  $\pm 10$  ppm window. Dynamic exclusion was set to place any selected  $m/z$  on an exclusion list for 45 seconds after a single MS/MS. All MS/MS spectra were searched against a protein database using Thermo Proteome Discoverer 1.3 (Thermo Fisher Scientific, San Jose, CA) using a *Rattus Norvegicus* protein database downloaded from Uniprot (<http://www.uniprot.org>). Fully tryptic peptides with up to 2 missed cleavage sites were selected. Variable modifications considered during the search included methionine oxidation (15.995 Da), and cysteine carbamidomethylation (57.021 Da). Proteins were identified at 99% confidence with XCorr score cut-offs (Qian et al., 2005) as determined by a reversed database search.

#### **4.2.6 Data analysis**

The protein and peptide identification results were visualized with Scaffold v 3.6.1 (Proteome Software Inc., Portland, OR, USA). Protein identifications with a minimum of two peptides identified at 0.1% peptide false discovery rate (FDR) were deemed correct. Significant changes in protein expression were identified by Fisher's exact test. Fold changes in protein expression were calculated using weighted normalized spectra with 0.5 imputation value.

The different comparisons were as follows:

- A) Control behavior verses  $\Delta 9$ -THC behavior
- B)  $\Delta 9$ -THC non-behavior verses  $\Delta 9$ -THC behavior
- C) Control non-behavior verses Control behavior



#### **4.2.7 IPA and gene ontology**

For pathway and ontology analysis, the protein list was filtered by applying the log<sub>2</sub>fc threshold of 0.6 and the Fisher's exact P-value of 0.05. Biological interpretation of identified proteins was performed by DAVID (Database of Annotation, Visualization and Integrated Discovery). The functional annotation clustering categorizes the differentially expressed proteins among the controlled vocabulary of Gene Ontology (GO) in three main categories: Biological process, which describes broad biological goal; Molecular function, which describes the task performed by individual gene products; and Cellular component, which describes the location of molecules inside the cell (Dennis et al., 2003). In current study, GO terms enriched with differentially expressed protein with a P-value  $\leq 0.05$  were selected.

Differentially expressed proteins were also analyzed in Ingenuity Pathway Analysis (IPA; Qiagen Inc., <https://www.qiagenbioinformatics.com>) to obtain canonical pathways, physiological functions, and networks associated with these proteins.

#### **4.2.8 Quantitative PCR (QPCR)**

Total RNA from the amygdala was extracted using the Qiagen RNeasy mini kit (Germantown, MD) and converted to cDNA by using the Qiagen RT2 cDNA synthesis kit provided with genomic DNA elimination buffer (Germantown, MD). Pre-designed gene specific primers, probes (Table 4.1) and Primetime Gene Expression Mastermix (Integrated DNA Technology, Coraville, IA) were used to perform real time QPCR. Each sample was run in triplicate in a final reaction volume of 20 ul according to manufacturer's instructions. The thermal cycler conditions were as follows: polymerase activation- 95C for 3 min, amplification- 40 cycles of 95C for 15 sec and 60C for 1 min.

In each experiment, standard curves for the gene of interest and the reference gene were made by using a range of cDNA dilutions (1:5-1:1000) and the PCR efficiency factor was calculated. The threshold Ct value of the gene of interest were normalized with Ct values of the reference gene, and PCR efficiency factor. Normalized Ct values were used to calculate the fold change from control by using method  $2^{(Ct(\text{Control mean}) - Ct(\text{Treatment}))}$  as described by Betancourt et al. (2007). Statistical analysis was conducted on the fold change values by using a non-parametric Kruskal-Wallis one way analysis of variance test to determine the significant treatment effect. The level of significance was set at  $p \leq 0.05$ .

Table 4.1 Primers, probes, and assay names for reference and target genes

Gene name	Assay name	Sequence 5'-3'
Actb	Rn.PT.	Primer 1 GGCATAGAGGTCTTTACGGATG
	39a.222214838.	Primer 2 TCACTATCGGCAATGAGCG
	g	Probe  /5HEX/TCCTGGGTA/ZEN/TGGAATCCTGTGGC/3IAkFQ
Gnai	Rn.PT.58.5104	Primer 1 GTCGTTTTCACTCTAGTTCTGAGA
	082	Primer 2 GAGTACCAGCTGAACGATTCCG
		Probe  /56FAM/CAACCAAAT/ZEN/TACATCCCAACCCAGCA3IABk  FQ

Table 4.1 (continued)

Gnas	Rn.PT.58.6088	Primer 1 CCTTCTCACCATCGCTGTT
	247	Primer 2 CACCATTGTGAAGCAGATGAG
		Probe
		/56FAM/CCGCCCTCT/ZEN/CCGTAAACCCATTAA/3IABkF
		Q
Gnaq	Rn.PT.58.3820	Primer 1 CACAAGCTCTTTATTGCGTCTA
	3772	Primer 2 CACTCTCAAGATCCCATAACAAGT
		Probe
		56FAM/ACACCTTCT/ZEN/CCACATCAACCTCTCGA/3IABkF
		Q
Gna1	Rn.PT.58.3405	Primer 1 GCCATCAACTTGTCTCCGT
3	4555	Primer 2 CACCATCTACAGCAACGTGA
		Probe
		56FAM/CTTCTCTCG/ZEN/GGCATCTACCAGCAC/3IABkF

### 4.3 Results and Discussion

In Chapter 2, we determined the effects of juvenile treatment with  $\Delta^9$ -THC on social behavior in adolescent rats. In Chapter 3, we determined the changes in amygdaloid proteomic expression and the associated canonical pathways in non-behavioral

adolescent rats treated with  $\Delta 9$ -THC as juveniles. Our lab has previously compared the changes in amygdaloid proteomic expression and the associated canonical pathways in control rats undergoing behavioral testing with non-behavioral control rats (Alugubelly et al., 2018). In this chapter, we extended those studies to determine the relationship between the changes in proteomic expression and the altered social behavior. This was done by comparing the differentially expressed proteins in the amygdala and the associated canonical pathways in (1) Control behavioral rats and  $\Delta 9$ -THC-treated behavioral rats and (2)  $\Delta 9$ -THC-treated behavioral rats and  $\Delta 9$ -THC-treated non-behavioral rats.

#### **4.3.1 Control behavioral verses $\Delta 9$ -THC-treated behavioral comparison**

In this analysis, protein validation using Scaffold identified 51 proteins that were differentially expressed. DAVID analysis revealed that GO term related to the biological process such as microtubule cytoskeleton organization and cell adhesion, and the cellular component such as cell leading edge (Figure 4.1). These GO terms are involved in axonal guidance (Dent and Gertler 2003), contact between axonal growth cone and target cell (Brose 1999) or neurite growth (Rehaine and Madox 2015).

Further analysis of these differentially expressed proteins using IPA revealed the enrichment of canonical pathways listed in table 1. Figure 4.2 indicates several important pathways based on the number of differentially expressed proteins involved, and the relevance of these pathways in terms of brain development. Some of these pathways are possibly related to CB1 signaling such as protein kinase A, CREB signaling, and cAMP signaling, phospholipase C and calcium signaling. Other pathways such as thrombin signaling and opioid signaling are important in terms of brain development or social play

(Figure 1B). Thrombin signaling, a key pathway involved in the neurite retraction in developing brains (Jalink 1994), and opioid signaling were inhibited in the  $\Delta 9$ -THC-treated behavioral rats as compared to the vehicle treated behavioral rats. Interestingly, opioid signaling was activated in  $\Delta 9$ -THC-treated non-behavioral rats compared to vehicle-treated behavioral rats, whereas it was inhibited in  $\Delta 9$ -THC-treated behavioral rats. This suggests that juvenile  $\Delta 9$ -THC exposure may have affected the opioids (receptors or peptides) that would cause altered binding and/or activation of opioid signaling during adolescence.

Thrombin signaling in neurons is activated when cells are going through several biological processes such as neurite retraction (Jalink et al., 1994), or apoptosis (Denovan et al., 1997). Other studies reported that thrombin is produced in the brain following the brain injury which may lead to break down of blood brain barrier (Xi et al., 2003). In our study, thrombin signaling was enriched in  $\Delta 9$ -THC-treated behavior rats compared to  $\Delta 9$ -THC-treated non-behavior rats but was not enriched in vehicle-treated behavior rats compared to vehicle-treated non-behavioral rats. This suggests that the juvenile  $\Delta 9$ -THC exposure caused some damage to the brain, including neurons in the amygdala causing them to synthesize thrombin. However, inhibition of thrombin signaling in  $\Delta 9$ -THC-treated behavior rats, may indicate an adaptive response elicited by amygdala neurons following social play.

In Chapter 3 we reported the mRNA expression of G protein subunits in non-behavior adolescent rats receiving  $\Delta 9$ -THC as juveniles. Because social play activates the CB1 and opioid signaling which share the common G protein mediated mechanism of activation (Trezza and Vanderschuren 2008), we were interested to measure the abundance of G

protein subunit mRNA in  $\Delta 9$ -THC-treated behavior rats. There was a 6% increase in the Gnai subunit, but Gaq mRNA was reduced by 13% relative to control (Figure 1C). Down regulation of Gaq may indicate the reversal effect of social play on signaling because in Chapter 3, we reported the up regulation of Gaq in  $\Delta 9$ -THC-treated non-behavior rats. Down regulation of Gaq mRNA may be associated with inhibition of calcium signaling and phospholipase C (PLC) signaling (Figure 4.2). In previous study, it was reported that CB1 agonist-mediated activation of PLC and  $Ca^{++}$  signaling occurs through binding of Gaq subunit to CB1 receptor (Lauckner et al., 2005). Since PLC is an enzyme involved in the biosynthesis of the CB1 ligand 2-arachidonoyl glycerol (2-AG; Kano et al., 2009), it can be hypothesized that developmental  $\Delta 9$ -THC exposure may cause long term alterations in 2-AG synthesis.

In previous studies, it has been demonstrated that CB1 signaling can have an enhancing effect on social play behavior (Trezza et al., 2008 a,b; Trezza et al., 2013; Manduce et al., 2015). However, there are also studies that report the opposite effect of CB1 activation on social play behavior (O'Shea et al., 2006; Schneider and Koch 2005). This disparity was addressed by Trezza and Vanderscuren (2008a,b) who, considering the high reward value of social play in rats, demonstrated that a cross-talk between cannabinoid and opioid receptors exists and this is responsible for the  $\Delta 9$ -THC-induced and/or CB1 activation-mediated enhancement of social play behavior. In the present study, we observed inhibition of opioid signaling in the  $\Delta 9$ -THC-treated rats when they were subjected to social play. A possible explanation may be the inhibition of related pathways such as CREB signaling and PKA signaling. Both of these pathways are inhibited due to down regulation of cAMP (Koh et al., 1997). CREB is a transcriptional

factor which is phosphorylated by PKA to its active form. The activation or inhibition of CREB in different brain regions can control different types of behaviors including anxiety and reward behaviors (Review: Carlezon et al., 2005). It has been demonstrated that microinjection of  $\Delta 9$ -THC into the amygdala inhibits CREB activation leading to an anxiogenic response in rats (Rubino et al., 2008). It was also reported that CREB activity is inhibited in the presence of opioid agonist, mainly because of the inhibitory action of opioid receptor on cAMP (Guitart et al., 1992). On the other hand, opioid antagonist activates CREB by releasing the tonic inhibition of opioid receptor on cAMP (Shaw-Lutchman et al., 2002). In present study, inhibition of cAMP, CREB, and PKA signaling in  $\Delta 9$ -THC treated rats is in agreement with previous studies.

Table 4.2 List of canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats receiving either corn oil or 10 mg/kg  $\Delta 9$ -THC during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play

Canonical pathways	Activity z-score
Protein Kinase A	Negative
Sirtuin signaling pathway	Negative
Thrombin signaling	Negative
CREB signaling in neurons	Negative
Calcium signaling	Negative
GM-SCF signaling	Negative
Ephrine receptor signaling	Negative
B cell receptor signaling	Negative
Integrin signaling	Positive
Phospholipase C	Negative
cAMP mediated signaling	Negative

Table 4.3 (continued)

iCOS-iCOSL signaling in T helper cells	Negative
Neuropathic pain signaling in dorsal horn neuron	Negative
Chemokine signaling	Negative
Synaptic long term potentiation	Negative
Glioma signaling	Negative
Melanin signaling	Negative
GNRH signaling	Negative
PKCB in T lymphocytes	Negative
PI3K in B lymphocytes	Negative

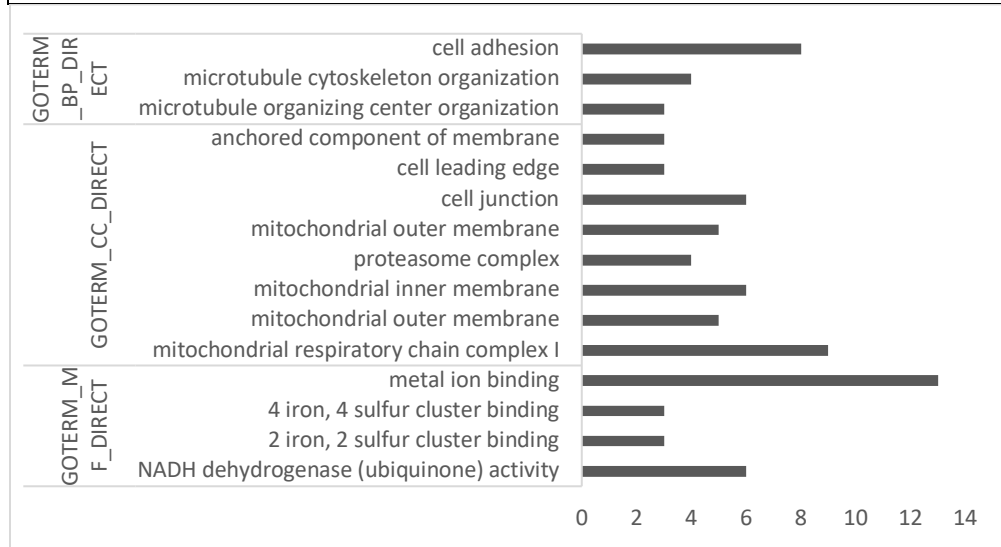


Figure 4.1 GO term analysis of differentially expressed proteins obtained from amygdala tissue of behavioral rats repeatedly exposed to either corn oil or 10 mg/kg  $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted 3-hours following social play on PND38.



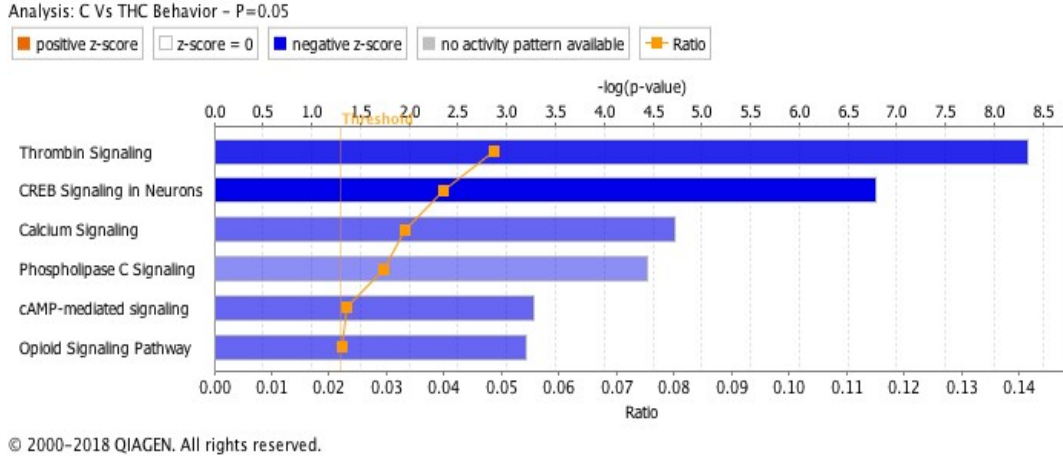


Figure 4.2 Relevant canonical pathways obtained from IPA analysis of differentially expressed proteins obtained from amygdala tissue of behavioral rats repeatedly exposed to either corn oil of 10 mg/kg  $\Delta$ 9-THC during juvenile ages (PND10-16). Tissue was extracted 3-hours following social play on PND 38.

Table 4.3 Differentially expressed proteins involved in the relevant canonical pathways shown in Figure 4.2.

Uniport Accession	Gene Symbol	Entrez Gene Name	Regulation
KCC4_RAT	CAMK4	Calcium/calmodulin dependent protein kinase IV	Down
KCC2A_RAT	CAMK2A	Calcium/calmodulin dependent protein kinase II	Down
FILUE2_RAT	CAMK2B	Calcium/calmodulin dependent protein kinase II beta	Down

Table 4.3 (continued)

KCC2D_RAT	CAMK2D	Calcium/calmodulin dependent protein kinase II	Down
A0A0G2K319_RAT	DUSP3	Dual specificity phosphatase	Down
GNA13_RAT	GNA13	G protein subunit alpha13	Down
M0R809_RAT	GNG4	G protein subunit gamma 4	Up
GRB2_RAT	GRB2	Growth factor receptor	Down
G3V9Y1_RAT	MYH10	Myosine heavy chain 10	Up
A0A0G2JSW0_RAT	MYL2A	Myosine light chain 12A	Up
PDE18_RAT	PDE18	Phosphodiesterase 18	Up
PPR18_RAT	PPP1R1B	Protein phosphatase 1 regulatory inhibition subunit	Up
RACK1_RAT	RACK1	Receptor for activated C kinase 1	Up



Figure 4.3 Relative change in mRNA expression of G protein subunits in amygdala of behavioral rats repeatedly exposed to either corn oil or 10 mg/kg of  $\Delta^9$ -THC during juvenile ages (PND10-16). Tissue was extracted 3-hours following social play on PND38.

#### 4.3.2 $\Delta^9$ -THC-treated behavioral versus $\Delta^9$ -THC-treated non-behavioral comparison

In this analysis, protein validation using Scaffold identified 64 proteins that were up regulated and 67 proteins that were down regulated. Functional clustering of these differentially expressed proteins was performed by using DAVID which identified GO term related to cellular component such as cell junction, cell-cell adherence and synapse (Figure 4.4).

Further analysis of these differentially expressed proteins using IPA revealed enrichment of canonical pathways listed in Table 4. The canonical pathways such as thrombin signaling and opioid signaling PKA, cAMP and CREB signaling are the same pathways that we observed to be altered in our previous experimental comparison discussed above. The differentially expressed proteins associated with these pathways are listed in table 5.

Additional pathways significantly enriched included dopamine degradation and dopamine DARPP32 feedback in cAMP signaling (Figure 4.5). The differentially expressed proteins involved in these pathways are listed in Table 5.

The dopamine degradation pathway involved only monoamine oxidase (MAO) A and B from our list of differentially expressed proteins. Down regulation of MAO may indicate the decrease in dopamine degradation as this enzyme is responsible for dopamine degradation. This finding is in agreement with previous studies that reported an increase in the neuronal firing in dopamine producing brain regions such as the ventral tegmental area and the substantia nigra (Renard et al., 2017; Wu et al., 2000) and an increase in the dopamine release (Oleson et al., 2012; Chen et al., 1990) following  $\Delta$ 9-THC exposure. However, an increase in the dopamine metabolites has also been observed following oral exposure to  $\Delta$ 9-THC (Novorro et al., 1993). This contradicts our findings in that we observed a down regulation of MAO which may not be correlated to increase in the dopamine metabolites. Dopaminergic neurotransmission is modulated by the retrograde action of CB1 receptors on GABA neurons which inhibits dopamine synthesis (Solinas et al., 2006).  $\Delta$ 9-THC-mediated activation of the CB1 receptors may have been responsible for  $\Delta$ 9-THC-mediated increase in dopamine levels observed by Novorro et al. (1993). It is also possible that the  $\Delta$ 9-THC-mediated increase in dopamine was due to the inhibitory action of  $\Delta$ 9-THC on MAO levels (Fisar 2010).

DARPP-32 is a protein phosphatase inhibitor mainly present in the dopaminergic neurons (Fernandez et al., 2006) and present in the brain regions which are enriched with dopamine innervations including the amygdala (Yger and Girault 2011). DARPP-32 is activated due to the stimulatory action of dopamine receptor (D1) on cAMP and PKA,

which phosphorylates DARPP-32 on Thr34 site. Thr34 phosphorylation eventually causes inhibition of CREB (Fernandez et al., 2006). Another study reported cyclin dependent kinase (cdk5) can phosphorylate DARPP-32 on Thr75 site which is responsible for inhibition of PKA (Bibb et al., 1999). Another study reported that DARPP-32 feedback in cAMP signaling pathway is activated when there is an interaction between opioid and dopaminergic neurotransmission (Mahajan et al., 2009). Also, the dual activation of opioid and dopamine receptor is associated with increase in the social play activity (Niesink and van Ree 1989). Therefore, inhibition of cAMP, CREB, PKA, opioid, dopamine degradation and DARPP-32 feedback signaling pathways found in our study may indicate that these pathways are inter related and play a role in mediating behaviors controlled by amygdala.

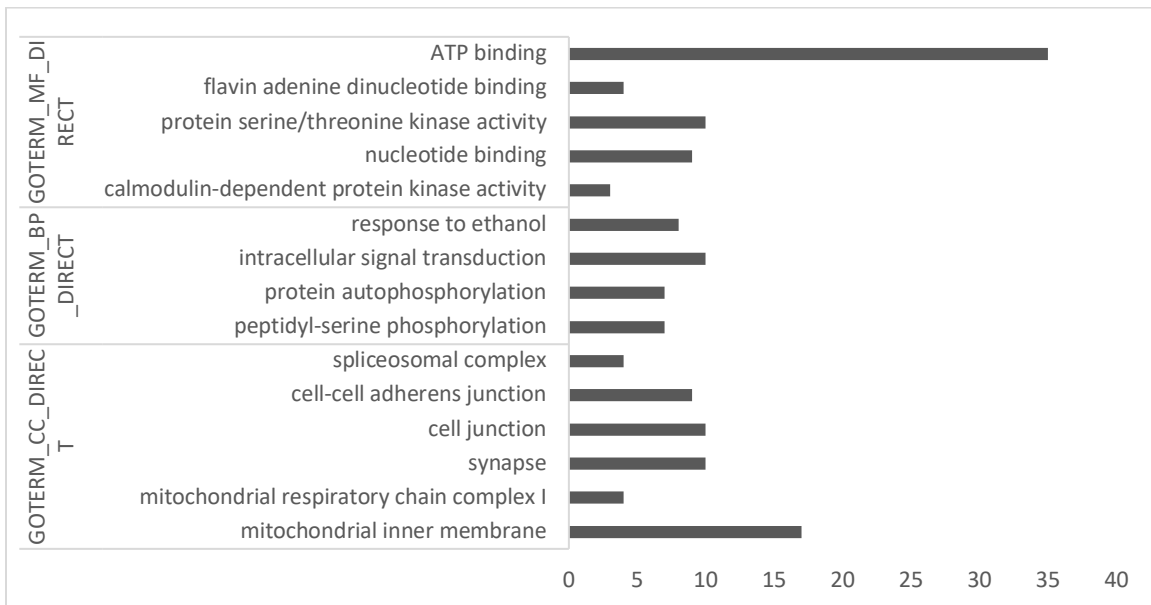


Figure 4.4 GO term analysis of differentially expressed proteins obtained from amygdala of adolescent rats repeatedly exposed to only  $\Delta 9$ -THC during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior  $\Delta 9$ -THC) or without being subjected to play (non-behavior  $\Delta 9$ -THC)

Table 4.4 List of canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats receiving only 10 mg/kg  $\Delta$ 9-THC during juvenile ages. Tissue was extracted on PND38, either 3-hours following play (behavior  $\Delta$ 9-THC) or without being subjected to play (non-behavior  $\Delta$ 9-THC).

Canonical pathways	Activity z-score
CREB signaling in neurons	Negative
Melatonin signaling	Negative
GNRH signaling	Negative
Thrombin signaling	Negative
Opioid signaling	Negative
Synaptic long term potentiation	Negative
CXCR4 signaling	Negative
CCR3 signaling in eosinophils	Negative
fMLP signaling in neutrophils	Negative
Androgen signaling	Negative
Chemokine signaling	Negative
Corticotrophin releasing hormone signaling	Negative
Protein kinase A signaling	Negative
GPCR signaling	Negative
Xenobiotic metabolism signaling	Negative
GPCR mediated nutrient sensing in enteroendocrine cells	Negative
IL-B signaling	Negative
UVC-induced MAPK signaling	Negative
Neuropathic pain signaling in dorsal horn neurons	Negative
Gaq signaling	Negative
G beta gamma signaling	Negative

Table 4.4 (continued)

Phospholipase C signaling	Negative
P2Y purigenic receptor signaling	Negative
UVB induced MAPK signaling	Negative
FC epsin RI signaling	Negative
Dopamine DRPP-32 feedback in cAMP signaling	Negative
Synaptic long term depression	Negative
GM-CSF signaling	Negative
Tryptophan degradation	Negative
Renin Angiotensin signaling	Negative
cAMP mediated signaling	Negative
NRF2-mediated oxidative stress	Negative
Neuroglin signaling	Negative
LPS-mediated MAPK	Negative
Ca induced T lymphocyte apoptosis	Negative
ErbB signaling	Negative
Cholestokinin mediated signaling	Negative
Thrombopiotin signaling	Negative
eNOS signaling	Negative
PPAR/RXR-alpha activation	Negative
PI3K signaling in B lymphocytes	Negative

Table 4.4 (continued)

Huntington disease signaling	Negative
CDK5 signaling	Negative
Endothelin signaling	Negative
GP6 signaling	Negative
Adrenomedulline signaling	Negative
P7056K signaling	Negative
Ca signaling	Negative
Aldosterone signaling in epithelial cells	Negative

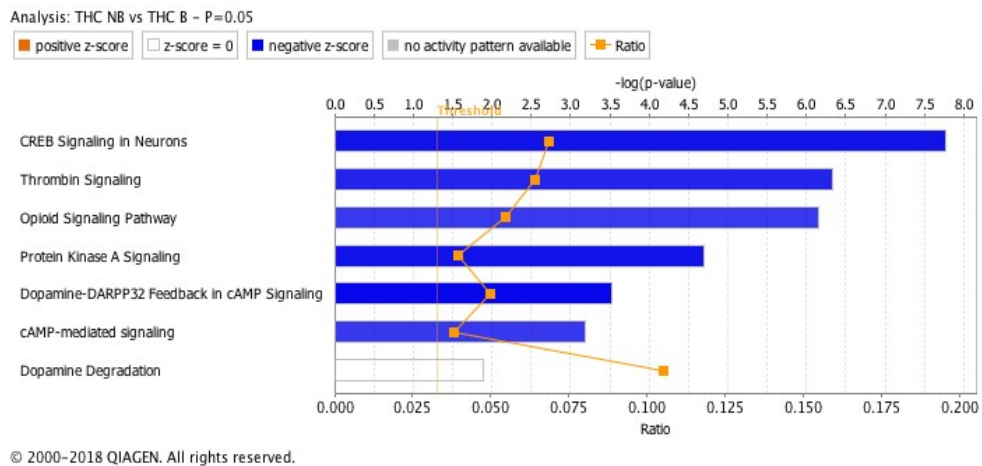


Figure 4.5 Relevant canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats receiving only 10 mg/kg  $\Delta^9$ -THC during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior  $\Delta^9$ -THC) or without being subjected to play (non-behavior  $\Delta^9$ -THC).



Table 4.5 Differentially expressed proteins involved in relevant canonical pathways shown in figure 4.5

Uniprot Accession	Gene Symbol	Entrez Gene Name	Regulation
FILN88_RAT	ALDH2	Aldehyde dehydrogenase	Down
G3V9G1_RAT	ADCYC	Adenylyl cyclase 5	Down
A0A0G2K7E5_RAT	CACNA2D1	Calcium voltage- gated channel auxiliary subunit alpha2delta 1	Down
KCC4_RAT	CAMK4	Calcium/calmoduline dependent protein kinase IV	Down
FILUE2_RAT	CAMk2B	Calcium/calmodulin dependent protein kinase II beta	Down
KCC2D_RAT	CAMK2D	Calcium/calmodulin dependent protein kinase II delta	Down
GNAS2_RAT	GNAS	GNAS complex locus	Up

Table 4.5 (continued)

M0R809_RAT	GNG4	G protein gamma subunit	Down
FILNE4_RAT	GRIA2	Glutamate ionotropic receptor AMPA subunit 2	Up
MAP2K1_RAT	MAP2K1	Mitogen activated protein kinase kinase 1	Down
FILP57_RAT	MAP2K4	Mitogen activated protein kinase kinase II	Up
KPCA_RAT	PRKCA	Protein kinase C alpha	Down
A0A0G2KSQ0_RAT	PRKCB	Protein kinase C beta	Down
KPCG_RAT	PRKCG	Protein kinase C gamma	Down
RACK1_RAT	RACK1	Receptor for activated C kinase	Up
S12AS_RAT	SLC12AS	Solute carrier family 12 member 5	Down

Table 4.5 (continued)

FILX13_RAT	PDE10A	Phosphodiesterase 10A	Down
CDK5_RAT	CDKS	Cyclin dependent kinase 5	Down
FILV89_RAT	RAPGAP	Rap1 GTPase	Down
AOFA_RAT	MAOA	Monoamine oxidase A	Down

### 4.3.3 Control behavior and control non-behavior comparison

In this analysis, protein validation using Scaffold identified 38 proteins that were up regulated and 64 proteins that were down regulated. Functional clustering of these differentially expressed proteins by using DAVID revealed that there was enrichment of molecular functions such as axonogenesis and positive regulation of axon elongation and enrichment of cellular components related to the synapse such as postsynaptic density and synaptic vesicles (Figure 4.6).

Further analysis of these differentially expressed proteins using IPA revealed enrichment of canonical pathways listed in table 6. Figure 3B indicates the most relevant pathways which are discussed in details. These pathways include opioid signaling, PKA signaling, cAMP signaling and dopamine degradation. Several studies reported the enhancing effect of opioid signaling on social play behavior (Trezza and Vanderschuren 2008 a,b; Vanderschuren et al., 1995 a,b; Beatty et al., 1982). As activation of opioid signaling in amygdala is mainly responsible for translating learned reward into motivation This

results in increased appetitive and/or consummatory behaviors towards that reward (Mahler and Berridge 2009). As play is a natural reward in rats, activation of opioid signaling in amygdala of behavioral rats was not surprising. The opioid signaling mainly occurs by binding of Gai subunits to opioid receptor causing inhibition of adenylyl cyclase enzyme. This causes decrease in cAMP and PKA signaling (Davis et al., 2003). Both cAMP and PKA signaling are decreased in our data.

The activation of opioid signaling was reversed in  $\Delta 9$ -THC-treated behavioral rats as discussed in the previous sections. This may indicate that juvenile  $\Delta 9$ -THC exposure altered opioid signaling in the brain that would alter the motivation for reward. However, despite the inhibition of opioid signaling,  $\Delta 9$ -THC-treated rats engaged in social play for more amount of time compared to controls as reported in chapter 2. Therefore, it can be hypothesized that juvenile  $\Delta 9$ -THC exposure causes functional changes in opioid signaling that would alter its activation during reward behaviors. Same pattern was observed for dopamine degradation pathway in both vehicle-treated and  $\Delta 9$ -THC-treated behavioral rats. In both comparisons, dopamine degradation pathway was enriched because of downregulation of MAO enzyme. This indicates inhibitory action of social play on dopamine degradation in amygdala. Previous studies have reported the enhancing effect of dopamine release on social play (Trezza et al., 2008; Peills et al., 1993; Neisink and van Ree 1989; Beatty et al., 1984). However, it is still need to be confirmed if the reduction in the MAO activity following social play can cause the accumulation of dopamine in amygdala and dopamine producing regions such as ventral tegmental area and striatum.

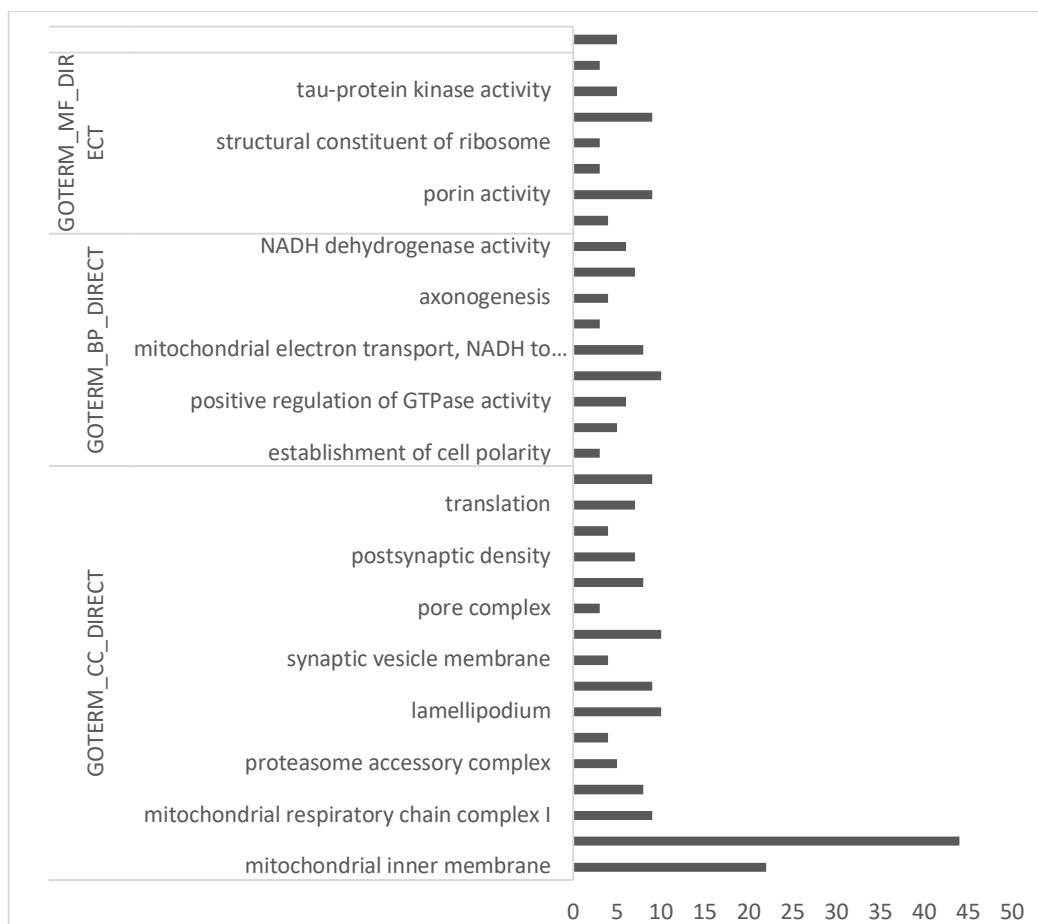


Figure 4.6 GO term analysis of differentially expressed proteins obtained from amygdala tissue of rats repeatedly exposed to corn oil during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior controls) or without being subjected to play (non-behavior controls).

Table 4.6 List of canonical pathways obtained from differentially expressed proteins in amygdala of adolescent rats exposed to corn oil during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior control) or without being subjected to play (non-behavior control).

Canonical pathways	Activity z-score
Mitochondrial dysfunction	No pattern
Sirbutin signaling	Positive

Table 4.6 (continued)

Glycogen degradation	Zero
Oxidative phosphorylation	Negative
Dopamine receptor signaling	Zero
Melatonin degradation	Zero
GDP-glucose biosynthesis	Zero
Glucose phosphatase degradation	Zero
Glycogen degradation	Zero
Protein kinase A signaling	Negative
Cardiac beta adrenergic signaling	Negative
Phenylalanine degradation	Zero
Serotonin receptor signaling	No pattern
Tryptophan degradation	Zero
Leukocyte extravasation	Zero
Dopamine degradation	Zero
Gai signaling	Zero
Noradrenaline degradation	Zero
Protein ubiquitnation	Zero
cAMP signaling	Negative
Relaxin signaling	Zero
Gaq signaling	Zero
Opioid signaling	Positive

Table 4.6 (continued)

NADH repair	Zero
tRNA splicing	Zero
Serotonin degradation	Zero
Melatonin degradation	Zero
GPCR signaling	Zero
EIF2 signaling	Zero
Gustation pathway	Zero
Semaphorin signaling	No pattern

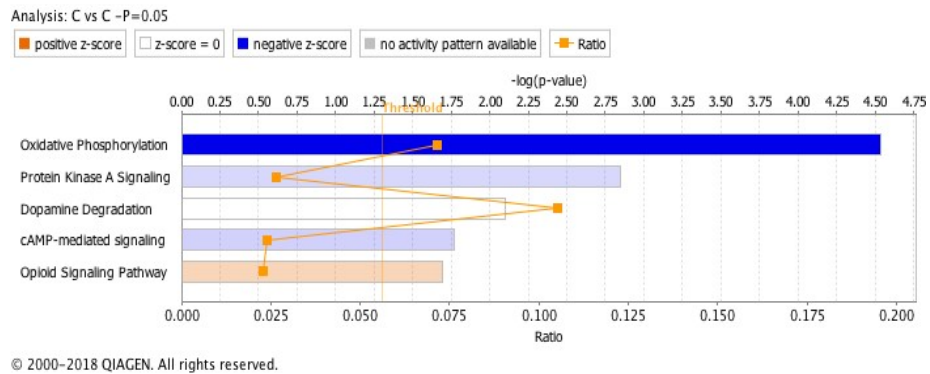


Figure 4.7 Relevant canonical pathways obtained from differentially expressed proteins obtained from amygdala of adolescent rats repeatedly exposed to only corn oil during juvenile ages (PND10-16). Tissue was extracted on PND38, either 3-hours following play (behavior controls) or without being subjected to play (non-behavior controls).

Table 4.7 List of differentially expressed proteins involved in canonical pathways listed in figure 4.

Uniprot Accession	Gene Symbol	Entrez Gene Name	Regulation
D3ZFQ8_RAT	CYC 1	Cytochrome c1	Down
NDUA9_RAT	NDUFA9	NADH Ubiquinone	Down
NDUAA_RAT	NDUAF10	NADH Ubiquinone	down
D4A0T0_RAT	NDUFB10	NADH Ubiquinone	Down
NDUS2_RAT	NDUSF2	NADH Ubiquinone	Down
UCR1_RAT	UQCRFS1	Ubiquinol cytochrome c	Down
G3V9G1_RAT	ADCYC5	Adenylyl cyclase 5	Down
AOFA_RAT	MAOA	Monoamine oxidase A	Down
AOFB_RAT	MAOB	Monoamine oxidase B	Down



Table 4.7 (continued)

A0A0G2KKB98_RAT	GSK3B	Glycogen synthetase kinase 3 beta	Up
FILX13_RAT	PDE10A	Phosphodiesterase 10A	Down
PDE1B_RAT	PDE1B	Phosphodiesterase 1B	Down
PP1R7_RAT	PPP1R7	Protein phosphatase 1 regulatory subnit 7	Down
RACK1_RAT	RACK1	Receptor for activated C kinase 1	Down
A0A0G2K5N6_RAT	ROCK2	Rho associated coiled coil containing protein	Up
FILV89_RAT	RAP1GAP	Rap1GTPase activating proteins	Down

Table 4.7 (continued)

D3ZWGZ_RAT	RGS7	Receptor for G protein signaling 7	Down
A0A0G2KB98_RAT	CLTA	Clathrin light chain A	Down

#### 4.4 Conclusion

This study evaluated the social play-mediated enrichment of canonical pathways in the amygdala and how these pathways are altered by developmental  $\Delta 9$ -THC exposure. We report that developmental  $\Delta 9$ -THC exposure alters normal opioid signaling during social play behavior as evidenced by the fact that engaging in social play by control rats activates opioid signaling, whereas prior exposure to  $\Delta 9$ -THC inhibits the social play-induced opioid signaling in rats. This is the most significant finding in this study because opioid signaling is mainly activated during the reward behaviors. We also found inhibition of thrombin signaling in  $\Delta 9$ -THC-treated behavioral rats. As thrombin signaling is activated in response to neuronal damage, enrichment of thrombin signaling in  $\Delta 9$ -THC treated rats may indicate a response of the neurons to the damage induced by developmental  $\Delta 9$ -THC exposure.

#### 4.5 Reference:

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## CHAPTER V

### CONCLUSION

There are numerous studies addressing the cognitive deficits in adults resulting from marijuana use during adolescence (Vadhan et al., 2017; Forti et al., 2014; Cass et al., 2014; Saez et al., 2014; Hajos et al., 2007) or gestation (Huizink 2014; Berghuis et al., 2007; Huizink et al., 2006). Given the increased popularity of ingesting marijuana edibles that has recently developed, this research investigated the persistent behavioral deficits and relevant signaling pathways that are altered following juvenile exposure to the psychoactive component of marijuana  $\Delta 9$ -THC.  $\Delta 9$ -THC is considered to be the main component responsible for producing the cognitive deficits associated with marijuana (D'Souza et al., 2005). The primary target of  $\Delta 9$ -THC in the brain is stimulation of the type I cannabinoid receptor (CB1) which is also involved in various aspects of brain development (Watson et al., 2008). Thus, the persistent neurological deficits associated with developmental marijuana exposure may be due to overstimulation of the CB1 receptor (Wilkinson et al., 2015) such that the normal progression of the brain development is disrupted.

Neurite outgrowth and axonal migration are two important steps in brain development that shape the connectivity of the brain by establishing proper synaptic connections across the regions (Fernandez-Ruiz et al., 2000). The CB1 receptor is consistently present in neurons undergoing these processes and is crucially important in the establishment of

the appropriate connections (Gaffari et al., 2012; Fernandez-Ruiz et al., 2000). During development, the acquisition of new behavioral skills depends upon the neuronal processes such as dendritic growth, the number of synapses and size of synapses in the brain (Dagwood et al., 1985). As juvenile period is associated with sharp increase in the number of synapses (Markaus and Petit 1987), over activation of CB1 receptor during this period can profoundly affect the acquisition of behavioral skills.

In our behavior data, we did not observe any direct effect of juvenile  $\Delta 9$ -THC exposure on anxiety in adolescent rats in the elevated plus maze (EPM). One possible reason could be the age at which the testing was conducted. Adolescence is a period characteristically linked with increased risk taking (Steinbern 2007). This natural tendency to take risks during adolescence may have simply increased the overall level of exploration and this decreased the ability to measure any increases or decreases in the level of anxiety during testing. We did observe significant differences in the social interaction test in adolescent rats exposed to  $\Delta 9$ -THC as juveniles. The most significant parameters affected involved increased levels of play behavior by the  $\Delta 9$ -THC-treated rats. A possible neurobiological explanation for this increased play behavior could be that the  $\Delta 9$ -THC exposure induced persistent changes that disrupted the inhibitory and excitatory balance in the brain. Activation of the CB1 receptor on the gamma-aminobutyric acid (GABA) neurons inhibits the release of GABA which reduces the inhibitory action of GABA in the brain. This 'dis-inhibition' has been reported to be enhanced by  $\Delta 9$ -THC treatment in rats (Radhakrishnan et al., 2015).

Our proteomic data identified the cellular pathways associated with  $\Delta 9$ -THC exposure and/or social play behavior in amygdala neurons. Thrombin signaling was the most

relevant pathway affected in  $\Delta 9$ -THC rats. Thrombin signaling in neurons is mainly responsible for growth cone collapse and neurite retraction (Jalink et al., 1994) which impairs the navigation of the axon and prevents it from reaching its target synapse (Harkany et al., 2008). In our data, enrichment of thrombin signaling was detected in  $\Delta 9$ -THC-treated behavioral rats and non-behavioral rats but was not detected in vehicle-treated rats which indicates that a long-term effect of repeated  $\Delta 9$ -THC ingestion during the juvenile period is altered thrombin signaling in neurons of the amygdala. As thrombin signaling in neurons is involved in neurite retraction (Jalink et al., 1994), is activated following brain injury (Vaughan et al., 1995) and is associated with neuronal death (Choi et al., 2003), the enrichment of thrombin signaling in  $\Delta 9$ -THC-treated rats in the current study may indicate that juvenile  $\Delta 9$ -THC exposure has altered one or more of these processes that thrombin signaling is linked to.

In addition, opioid signaling was enhanced in  $\Delta 9$ -THC-treated non-behavioral rats as compared to control non-behavioral rats but was inhibited in  $\Delta 9$ -THC-treated behavioral rats as compared to control behavioral rats. The different effects of  $\Delta 9$ -THC exposure on opioid signaling induced by the presence or absence of social behavior may suggest altered competitive binding of the cannabinoid and/or opioid receptors to G protein subunits. The cannabinoid and opioid receptors share a common G protein subunit and dual activation of both cannabinoid and opioid receptors during social play (Trezza and Vanderschuren 2008) in  $\Delta 9$ -THC-treated rats may have attenuated the binding of Gai subunit to opioid receptor as previously reported (Rios et al., 2006). Since opioid signaling is mainly associated with nociception and repeated  $\Delta 9$ -THC treatment reduces the response to painful stimuli even long after treatment ceases (Martin et al., 2004;

Caroline et al., 2000), this suggests that repeated  $\Delta 9$ -THC dampens the response of opioid signaling to stimuli. Our results agree with this premise in that introduction of a behavioral challenge resulted in the inhibition of opioid signaling in the amygdala of rats treated with  $\Delta 9$ -THC.

Overall, this study investigated the persistent effects of oral  $\Delta 9$ -THC exposure during the juvenile period and included the behavioral alterations observed and the cellular pathways that were either activated or inhibited in the amygdala of those exposed rats once they reached adolescence. This research has limitations in that it only tested the persistent alterations in behaviors related to anxiety and social interactions in response to exposure to  $\Delta 9$ -THC. First, our EPM data suggest an increase in locomotor activity in  $\Delta 9$ -THC-treated rats. Future studies should focus on validating the effects on  $\Delta 9$ -THC on locomotor activity using a more appropriate behavioral test specific for measuring this type of behavior. Secondly, we observed  $\Delta 9$ -THC-mediated effects on thrombin signaling in the amygdala neurons. Both thrombin signaling (Harkany et al., 2008) and the CB1 receptor (Fernandez-Ruiz et al., 2000) are involved in the migration of axons to target neurons in order to form a synapse. Therefore, future studies should also include investigation of the potential alterations in synaptic density throughout the entire brain following oral juvenile exposure to  $\Delta 9$ -THC. Thirdly, we observed that in  $\Delta 9$ -THC-treated behavior rats, there is an alteration in opioid signaling. This may suggest alterations in the binding affinity of G protein subunits to the opioid receptor in the amygdala. Future studies should investigate the binding affinity of G protein subunits to the opioid receptor in animals previously treated with  $\Delta 9$ -THC by administering a challenge that would activate opioid receptors.

While the popularity of edible marijuana products is increasing in the United States, the data obtained in our study may provide an insight to the increasing popularity of marijuana and related products in the United States. Our data has clearly established a link between repeated juvenile  $\Delta 9$ -THC exposure and deficits in social behaviors during adolescence. This effect of  $\Delta 9$ -THC on the developing brain in juveniles may challenge to the public's attitude to support the popularity of marijuana products. As addressed by previous studies, the child friendly packaging of marijuana edibles and the lack of limitations on the maximum content of  $\Delta 9$ -THC in a single edible are major threats to the children. Our data suggest that consumption of edibles by juveniles may be catastrophic in terms of the acquisition of skills which are necessary for meeting the challenges in future life. The results of this study also strongly suggest that care should be taken when popularizing marijuana edible products in the media so that children will not misinterpret marijuana as being safe for them to ingest.

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## APPENDIX A

### Lists of Proteins Obtained from Proteomic Analysis Under Different Experimental Conditions

Table A.1 List of differentially expressed proteins obtained from vehicle or  $\Delta^9$ -THC treated non-behavior rats

Uniprot Accession	Fishers exact P-value	LOG2FC
TBA3_RAT	0.015	0.378511623
G3P_RAT	0.022	0.584962501
MBP_RAT	0.029	0.584962501
ATPB_RAT	0.035	0.485426827
ACON_RAT	0.072	0.485426827
ALBU_RAT	0.019	-0.152003093
DYN1_RAT	0.048	0.485426827
1433Z_RAT	0.065	0.485426827
F1MA36_RAT	0.0019	-0.321928095
VATB2_RAT	0.047	0.485426827
Q6T487_RAT	0.0015	-0.514573173
KCC2A_RAT	0.0035	0.765534746
DPYL4_RAT	0.04	0.678071905
CALR_RAT	0.059	-0.321928095
G3V741_RAT	0.039	-0.321928095
D4A0F5_RAT	0.059	-0.321928095
A0A0G2K5Q0_RAT	0.047	0.765534746
G3V9Y1_RAT	0.022	-0.514573173
Q6IMZ3_RAT	0.0055	1.201633861
DCE2_RAT	0.026	1
D3Zfq8_RAT	0.039	-0.514573173
PP1R7_RAT	0.064	-0.514573173
F1LX07_RAT	0.061	-0.514573173
ROA1_RAT	0.056	-0.514573173
F1LMH0_RAT	0.057	-0.736965594
F1LN88_RAT	0.072	1.070389328
HPCL4_RAT	0.017	-0.736965594
A0A0G2K079_RAT	0.032	1.485426827
NDUAA_RAT	0.023	-1
A0A0A0MY13_RAT	0.032	1.378511623
RHOB_RAT	0.043	1
F1LPG3_RAT	0.0018	2.744161096
Q5RK08_RAT	0.061	1.263034406
WASF1_RAT	0.061	1.263034406

Table A.1 (continued)

A0A0G2JUT1_RAT	0.066	-0.736965594
RACK1_RAT	0.0039	-1.321928095
D3ZFX4_RAT	0.011	4.087462841
ADHX_RAT	0.038	-1
LIN7A_RAT	0.035	1.925999419
DC1I1_RAT	0.071	1.632268215
B0BNL2_RAT	0.038	-1.321928095
PRS6B_RAT	0.022	2.201633861
REEP5_RAT	0.0046	-1.736965594
D4AB17_RAT	0.0092	2.137503524
PENK_RAT	0.032	-1
DC1L1_RAT	0.0099	-2.321928095
F1M1Y0_RAT	0.035	2.070389328
USO1_RAT	0.0014	-4.321928095
A0A0G2K459_RAT	0.078	-3.321928095
F1M7X3_RAT	0.0099	-2.321928095
G3V8K2_RAT	0.066	-1
G3V8R2_RAT	0.0053	-2.321928095
YKT6_RAT	0.033	3.807354922
B4F7C7_RAT	0.0056	-3.836501268
G3V940_RAT	0.064	1.432959407
Q5EBD0_RAT	0.019	-1.321928095
6PGL_RAT	0.022	2.201633861
PSB3_RAT	0.021	-3.473931188
A0A0G2K3V4_RAT	0.071	-1
PSB1_RAT	0.0083	2.432959407
A0A0G2KA27_RAT	0.035	2.070389328
F1M013_RAT	0.041	-3.321928095
F1LM19_RAT	0.038	-1.321928095
TMM35_RAT	0.078	-3.321928095
A0A0G2K5N6_RAT	0.019	4
A0A0G2K6T9_RAT	0.018	-1.736965594
Q6P3V9_RAT	0.0099	-1.736965594
SFXN5_RAT	0.025	-1.321928095
PPM1E_RAT	0.021	-3.64385619
UBA5_RAT	0.058	-1.736965594

Table A.1 (continued)

AGAP2_RAT	0.033	3.700439718
A0A0G2JY03_RAT	0.033	-1.736965594
RCN2_RAT	0.0051	2.5360529
Q5RJN0_RAT	0.043	-1.321928095
RD23B_RAT	0.033	-1.736965594
MPI_RAT	0.078	-3.321928095
NTF2_RAT	0.033	-1.736965594
PRI0_RAT	0.041	-3.321928095
PSMD1_RAT	0.056	1.925999419
D4A5X7_RAT	0.058	-1.321928095
PHYIP_RAT	0.023	-1.321928095
D3ZT36_RAT	0.033	3.700439718
ZWINT_RAT	0.078	-3.321928095
CHRD1_RAT	0.033	3.700439718
AL7A1_RAT	0.055	3.584962501
TXTP_RAT	0.033	-1.736965594
G3V7Q6_RAT	0.055	3.584962501
G3V7U4_RAT	0.071	-1.321928095
G3V7I8_RAT	0.055	3.584962501
O88321_RAT	0.055	3.584962501
PSB2_RAT	0.078	-3.321928095
D3ZRM9_RAT	0.041	-3.473931188
M0RC57_RAT	0.011	4.087462841
A0A0G2JT00_RAT	0.055	3.584962501
sp P62243 RS8_RAT	0.041	-3.321928095
G3V914_RAT	0.043	-1.736965594
G3V912_RAT	0.033	3.700439718
PA2G4_RAT	0.011	-3.836501268
A0A0G2K724_RAT	0.021	-3.473931188
B2RYS8_RAT	0.078	-3.321928095
RLA1_RAT	0.078	-3.321928095
RUVB1_RAT	0.055	3.584962501
B1H2A2_RAT	0.078	-3.321928095
RLA0_RAT	0.021	-3.473931188
D3Z9G9_RAT	0.041	-3.321928095
D3ZM33_RAT	0.078	-3.321928095

Table A.1 (continued)

A0A0G2K526_RAT	0.056	2.201633861
M0R7P0_RAT	0.078	-3.321928095
Q505I9_RAT	0.078	-3.321928095

Table A.2 List of differentially expressed proteins obtained from vehicle or  $\Delta$ 9-THC treated behavior rats

Uniprot Accession	Fishers Exact P-Value	LOG2FC
KCC2A_RAT	0.001	0.736965594
D4A8U7_RAT	0.00074	-1
F1LUE2_RAT [5]	0.0074	0.736965594
NDUS1_RAT	0.034	0.584962501
Q5XIH3_RAT	0.0082	1
D3ZA84_RAT	0.014	-1
CALX_RAT	0.042	0.678071905
A0A0G2K9J2_RAT	0.018	-1
KCC2D_RAT	0.013	0.736965594
G3V9Y1_RAT	0.0091	0.925999419
VDAC2_RAT	0.0032	1.070389328
NFM_RAT	0.0076	1.201633861
LSAMP_RAT	0.039	-1
A0A0G2JZM8_RAT	0.025	-1
UCRI_RAT	0.023	1.137503524
Q5M7T6_RAT	0.00042	1.736965594
PALM_RAT	0.0093	1.378511623
RACK1_RAT	0.023	1.137503524
PPR1B_RAT	0.02	1.5360529
GRB2_RAT	0.04	1.736965594
A0A0G2K8V2_RAT (+2)	0.046	1.321928095
PDE1B_RAT	0.011	2.201633861
CALB1_RAT	0.0068	1.632268215
A0A0G2JSW0_RAT [2]	0.044	1.137503524
A0A0G2K261_RAT	0.034	3.584962501
ACYP2_RAT	0.034	3.584962501
A0A0G2K3I9_RAT	0.012	2.321928095
RAP2B_RAT [2]	0.02	1.5360529
G3V7L6_RAT (+1)	0.03	1.736965594

Table A.2 (continued)

FUBP1_RAT	0.01	3.906890596
NDUV2_RAT	0.025	1.378511623
ERR1_RAT	0.0018	4.321928095
I6L9G6_RAT	0.033	1.321928095
AGAP2_RAT	0.036	3.473931188
G3V7Q6_RAT	0.019	3.700439718
NDUS2_RAT	0.0056	4.087462841
KCC4_RAT	0.02	1.736965594
ACBP_RAT [2]	0.02	3.836501268
A0A0H2UI10_RAT	0.036	-3.64385619
B2GUY4_RAT (+1)	0.034	3.584962501
A0A0G2K0P8_RAT	0.034	3.584962501
Q6PDW1_RAT	0.036	3.473931188
D3Z955_RAT	0.02	3.836501268
TOM34_RAT	0.036	-3.64385619
sp K22E_HUMAN	0.0014	2.321928095
M0R809_RAT	0.034	3.584962501
G3V964_RAT	0.0061	4.058893689
M0R4B6_RAT (+1)	0.018	2.070389328
A0A0G2KAZ7_RAT	0.019	3.700439718
GNA13_RAT	0.03	1.736965594
G3V8H3_RAT	0.036	-3.64385619

Table A.3 List of differentially expressed proteins obtained from  $\Delta$ 9-THC treated behavior or non-behavior rats

Uniprot Accession	Fishers Exact P-Values	LOG2FC
D4ABN3_RAT	0.0043	0.678071905
GPM6A_RAT	0.0017	0.925999419
D3ZDU5_RAT	0.03	0.765534746
PCLO_RAT	0.038	1
PGRC1_RAT	0.016	0.925999419
NFM_RAT	0.0082	1.263034406
A0A0G2JZH8_RAT	0.0087	1.321928095
A0A0G2JUT1_RAT	0.049	0.925999419
PCYOX_RAT	0.0052	1.5360529
F1LRZ7_RAT	0.041	2.03562391
D4A435_RAT	0.021	1.722466024
CX6C2_RAT	0.024	3.700439718

Table A.3 (continued)

THIC_RAT	0.002	2.560714954
A0A0G2KAW4_RAT	0.0052	2.232660757
RACK1_RAT	0.013	1.321928095
B4F774_RAT	0.021	1.632268215
CSKI1_RAT	0.025	1.378511623
F1LV13_RAT	0.041	1.5360529
B0BNL2_RAT	0.014	1.5360529
ELOB_RAT	0.014	3.906890596
RTCB_RAT	0.014	3.906890596
A0A0G2K926_RAT	0.0004	4.584962501
sp P62909 RS3_RAT	0.014	3.906890596
B1WC34_RAT	0.0055	2
F1LP57_RAT	0.015	1.847996907
PSA2_RAT	0.024	1.584962501
Q5EBD0_RAT	0.014	1.5360529
CDS2_RAT	0.0089	2.10433666
D3ZWA8_RAT	0.0052	2.201633861
SCAM1_RAT	0.025	2.292781749
D4A781_RAT	0.0044	4.169925001
PPM1E_RAT	0.0013	4.392317423
LASP1_RAT	0.041	1.722466024
B4F7C7_RAT	0.0078	4
A0A0G2K6T9_RAT	0.0052	2.232660757
G3V6L9_RAT	0.014	3.906890596
DNJA2_RAT	0.0089	2.10433666
M0R735_RAT	0.0024	4.247927513
A0A0G2K0V8_RAT	0.024	3.700439718
A0A0H2UHF6_RAT	0.025	1.847996907
D3ZDC0_RAT	0.0044	4.169925001
CSRP1_RAT	0.014	3.906890596
ZWINT_RAT	0.043	3.584962501
ACYP2_RAT	0.043	3.584962501
A0A0G2JZB8_RAT	0.014	3.906890596
A0A0G2K7W7_RAT	0.024	3.700439718
FUBP1_RAT	0.014	3.807354922
Q6P3V9_RAT	0.025	1.722466024
SCN2B_RAT	0.024	3.700439718
MPI_RAT	0.024	3.700439718
A0A0G2K0P8_RAT	0.043	3.584962501
PRRT2_RAT	0.014	3.906890596
UK114_RAT	0.041	2.137503524
LYPA2_RAT	0.043	3.459431619

Table A.3 (continued)

SGT1_RAT	0.043	3.459431619
PRIORAT	0.014	3.906890596
F1LNE4_RAT	0.037	1.378511623
NPTX1_RAT	0.024	3.700439718
AP1M1_RAT	0.043	3.459431619
M0R907_RAT	0.043	3.459431619
sp P21531 RL3_RAT	0.0078	4
A0A0G2JVS2_RAT	0.043	3.459431619
M0R809_RAT	0.043	3.459431619
RAB4B_RAT	0.00022	4.700439718

Table A.4 List of differentially expressed proteins obtained from  $\Delta$ 9-THC treated behavior or non-behavior rats

Uniprot Accession	Fishers Exact P-Values	LOG2FC
F1LNK0_RAT	0.033	0.584962501
M0R8M9_RAT	0.022	0.584962501
D4A8U7_RAT	0.0016	1.263034406
Q499Q4_RAT	0.036	1.201633861
RUFY3_RAT	0.048	1.070389328
ERR1_RAT	0.0056	4.321928095
G3V6L4_RAT	0.029	1.5360529
D3ZHY9_RAT	0.0051	1.765534746
A0A0G2K4X8_RAT	0.02	1.584962501
A0A0G2JSZ5_RAT	0.024	1.432959407
PACS1_RAT	0.0077	2.5360529
Q5U2P5_RAT	0.048	1.807354922
I6L9G6_RAT	0.047	1.722466024
Q5M9H7_RAT	0.032	1.847996907
G3V6T1_RAT	0.047	2.070389328
LASP1_RAT	0.0094	4.247927513
PCYOX_RAT	0.0038	2.40599236
Q5U2S7_RAT	0.047	1.925999419
PRS7_RAT	0.03	2.201633861
A0A0G2KB98_RAT	0.016	4.087462841
ELOB_RAT	0.047	1.925999419
RTCB_RAT	0.042	3.700439718
LYAG_RAT	0.016	4.087462841



Table A.4 (continued)

M0R735_RAT	0.047	2.070389328
A0A0G2K5N6_RAT	0.0056	4.321928095
A0A0G2K7W7_RAT	0.032	1.925999419
CSRP1_RAT	0.042	3.700439718
PSMD1_RAT	0.047	2.070389328
O88321_RAT	0.042	3.700439718
PSB1_RAT	0.047	2.070389328
B5DFN4_RAT	0.042	3.807354922
D3ZWA8_RAT	0.026	3.906890596
PPM1H_RAT	0.042	3.807354922
RCN2_RAT	0.03	2.201633861
D3ZPR0_RAT	0.042	3.700439718
D3Z955_RAT	0.042	3.807354922
A0A0G2JT00_RAT	0.026	3.906890596
D3Z981_RAT	0.016	4.087462841