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Reelin deficiency contributes to long-term behavioral abnormalities induced by chronic adolescent exposure to 9-tetrahydrocannabinol in mice

Attilio Iemolo¹, Patricia Montilla-Perez¹, Jacques Nguyen², Victoria B Risbrough^{2,3}, Michael A Taffe², Francesca Telese¹

¹Department of Medicine, University of California, San Diego, La Jolla, CA 92093, USA

²Department of Psychiatry, University of California, San Diego, La Jolla, CA 92093, USA

³Center for Excellence in Stress and Mental Health, Veterans Affairs San Diego Healthcare System, San Diego, CA 92161, USA

Abstract

Cannabis use is widespread among adolescents and has been associated with long-term negative outcomes on neurocognitive functions. However, the factors that contribute to the long-term detrimental effects of cannabis use remain poorly understood. Here, we studied how *Reelin* deficiency influences the behavior of mice exposed to cannabis during adolescence. *Reelin* is a gene implicated in the development of the brain and of psychiatric disorders. To this aim, heterozygous Reeler (HR) mice, that express reduced level of *Reelin*, were chronically injected during adolescence with high doses (10mg/kg) of 9-tetrahydrocannabinol (THC), a major psychoactive component of cannabis. Two weeks after the last injection of THC, mice were tested with multiple behavioral assays, including working memory, social interaction, locomotor activity, anxiety-like responses, stress reactivity, and pre-pulse inhibition. Compared to wild-type (WT), HR mice treated with THC showed impaired social behaviors, elevated disinhibitory phenotypes and increased reactivity to aversive situations, in a sex-specific manner. Overall, these findings show that *Reelin* deficiency influences behavioral abnormalities caused by heavy consumption of THC during adolescence and suggest that elucidating Reelin signaling will improve our understanding of neurobiological mechanisms underlying behavioral traits relevant to the development of psychiatric conditions.

* Correspondence should be addressed to ftelese@ucsd.edu.

CRedit authorship contribution statement

Attilio Iemolo: Conceptualization, Investigation, Methodology, Writing- Original Draft; **Patricia Montilla-Perez:** Validation; **Jacques Nguyen:** Investigation for measurements of THC plasma levels; **Victoria B Risbrough:** Conceptualization for the PPI experiments; **Michael A Taffe:** Supervision for measurements of THC plasma levels; **Francesca Telese:** Conceptualization, Formal analysis, Supervision, Writing- Review & Editing.

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1. Introduction

Heavy and frequent cannabis use by adolescents has been linked epidemiologically to increased risk of developing psychiatric conditions, including schizophrenia, psychosis, and substance use disorders (Volkow, 2016). Similarly, animal studies show that administration of cannabinoids (e.g. THC), during adolescence, perturbs a wide range of behaviors, including memory, social interaction, anxiety, and sensorimotor gating by targeting the endocannabinoid (eCB) system (Rubino et al., 2015). Despite the evidence of possible detrimental health outcomes associated with adolescent cannabis use, a recent survey in the US revealed a substantial increase in daily use of cannabis and a decreased perception of the risks associated with its regular use by adolescents (Johnston, 2018). Further, the increasing legalization of recreational cannabis use has led to calls to understand whether such policies put adolescents at higher risk of developing psychiatric disorders. All this emphasizes the need to better understand the neurobiological mechanisms associated with heavy consumption on cannabis during adolescence (Wilkinson et al., 2016).

During adolescence, the brain undergoes continuous remodeling of its structure, connectivity, and plasticity (Sturman and Moghaddam, 2011; Arain et al., 2013). In addition, substantial hormonal changes during adolescence influence not only reproductive functions, but also the emergence of sex differences in cognitive, social, and emotional behaviors (Schulz and Sisk, 2016). Thus, the adolescence is considered a critical period wherein brain development may be altered by the exposure to psychoactive drugs, which can lead to sex-specific behavioral abnormalities and increased risk for psychopathology in adulthood (Cousijn et al., 2018; Lisdahl et al., 2018).

However, the factors that contribute to long-term detrimental effects of cannabis exposure during adolescence remain poorly understood. The goal of this study is to examine the potential role of Reelin signaling in modulating the behavioral effects of cannabis on the adolescent brain.

Reelin is a protein of the extracellular matrix that is predominately expressed in neuronal cells and plays a key role in brain development and synaptic plasticity (D'Arcangelo et al., 1995). During embryonic stages, Reelin activates an extensive signaling cascade that is critical for the proper migration and cell positioning of cortical neurons (Sekine et al., 2014). During adolescence, Reelin signaling promotes the development of the synaptic excitation/inhibition (E/I) balance within the prefrontal cortex (Iafrati et al., 2014; Bouamrane et al., 2016). In the mature brain, Reelin is required for learning and memory by regulating the N-methyl-D-aspartate receptors (NMDA-R) function and the expression of neuronal activity-dependent genes (Weeber et al., 2002; Qiu et al., 2006; Niu et al., 2008; Rogers et al., 2011; Telese et al., 2015). In humans, *Reelin* deficiency has been linked to the development of psychiatric disorders (Ishii et al., 2016). Thus, the HR mice, that expressed lower level of Reelin, have been proposed as a valid animal model to study neurodevelopmental psychiatric disorders (Lossi et al., 2019). Whether there is a functional relationship between *Reelin* deficiency and the consequences of adolescent exposure to high levels of THC remains unknown.

To this aim, we examined the long-lasting behavioral outcomes of chronic adolescent exposure to high doses of THC (10mg/kg) in female and male HR mice. We compared HR mice to their WT littermate controls in a battery of behavioral tests exploring different facets of cognitive and emotional responsiveness, including working memory (Sannino et al., 2012), social interaction (Yang et al., 2011), anxiety-like responses (Bailey and Crawley, 2009), stress reactivity (Can et al., 2012), and pre-pulse inhibition (Geyer et al., 2002). This is the first study to investigate the relationship between *Reelin* deficiency and the effect of adolescent exposure to THC in mice.

2. Materials and Methods

2.1 Animals.

All experimental procedures were approved by the institutional animal care and use committee at University of California, San Diego. Mice were housed (3–4 per cage) under a 12h light/12h dark cycle and provided with food and water ad libitum. HR mice were bred in house using the B6C3Fe a/a-ReInrl/J line (The Jackson Laboratory, #000235) (D'Arcangelo et al., 1995).

2.2. Drug treatment protocol and experimental design for behavioral analysis.

THC was provided by the U.S. National Institute on Drug Abuse and was dissolved in a vehicle solution consisting of ethanol, tween, and 0.9% saline (1:1:18) on the day of administration. The “high” dose and chronic treatment protocol were selected to study the behavioral effects of heavy and chronic cannabis exposure. The dosage we used (10 mg/kg) was referred to as ‘high’ based on previous studies using low to high ranges of THC in mice (Trexler et al., 2018; Kasten et al., 2019). This dose was also selected to achieve physiologically relevant amounts of THC in mice based on plasma level concentrations observed in cannabis use in humans, as shown in Suppl. Fig. 1 and further explained in the discussion (Huestis et al., 1992; Abrams et al., 2007; Zuurman et al., 2008; Karschner et al., 2009; Nguyen et al., 2016). Vehicle or THC were administered daily to adolescent mice by intraperitoneal injections from post-natal day (PND) 28 to PND 48, which cover the adolescent period in mice (Laviola et al., 2003) (Fig. 1A). To examine the long-term effects of chronic adolescent exposure to THC, mice were tested two weeks after a drug abstinence period, starting at PND 63. We used 9 cohorts of mice to perform multiple behavioral assays. Cohorts 1 to 5 were subjected to locomotor activity, open field (OF), six-different objects (6-DOT), light-dark (LD), three-chamber social approach, and tail suspension (TS) tests. Cohorts 6 to 9 were subjected to locomotor activity, acoustic startle response (ASR) and pre-pulse inhibition (PPI) tests. Mice were tested between 10:00 am and 5:00 pm. Behavioral assays were conducted on separate days and all behavioral tests were performed once on each mouse. The number of mice used in each behavioral assay is reported in Table 1. To prevent bias due to olfactory cues, the behavioral apparatus was cleaned with diluted ethanol solution in between mice.

2.3. Six different objects test.

We used the 6-DOT to study short-term working memory in mice, as previously described (Sannino et al., 2012). First, mice were left free to explore an empty arena (60×40×35 cm)

during a 10 min habituation trial. Afterwards, mice were exposed to six different objects for a total time of 10 min (familiarization trial). After an inter-trial interval of 1 minute, mice were exposed to identical copies of the familiar objects, but one object was substituted with a novel object (test trial). The exploration time across different trials was measured with Anymaze (Ugo Basile, Varese, Italy) and was used to calculate a discrimination index (DI) as follows: (time spent exploring novel object – average time spent exploring familiar objects) / Total time spent exploring novel + familiar objects. The time exploring the novel object and the DI were used as indexes of novelty-induced exploratory activity and working memory, respectively.

2.4. Three-chamber social approach test.

The three-chamber social approach test was used to measure social behaviors, as previously described (Yang et al., 2011). The apparatus comprised of three-chambered box with dividing walls with small openings to allow free exploration of the three chambers, such as one empty central chamber, one side chamber containing an empty small wire cage (novel object) and one side chamber containing a stranger mouse inside a small wire cage (novel mouse). The target mouse was first placed in the center chamber and allowed to explore the apparatus for 15 min. After introducing the novel mouse and the novel object in the side chambers, the mouse was allowed to explore for 10 min. The placement of the novel mouse or novel object in the left or right chambers was systematically alternated in between trials. The time spent in each compartment and the time spent actively sniffing the novel mouse or the novel object were manually scored. Longer time spent with or exploring the novel mouse versus the novel object was considered an index of sociability. Sociability index (SI) was calculated as [(time spent exploring or sniffing novel mouse - time spent exploring or sniffing novel object) / (total time spent exploring or sniffing novel mouse and novel object)].

2.5. Tail suspension test.

The TS test was used to measure motor responses under aversive conditions (Can et al., 2012). Mice were suspended by their tails with tape to a bar in a position that they could not escape or hold on to nearby surfaces. The test lasted for 6 minutes and the mobility time (s) was manually scored for each minute. We analyzed was the mobility time (s) as the sum of the final 5 minutes.

2.6. Open field test.

The OF test was used to measure anxiety-like behavior (Bailey and Crawley, 2009). Mice were placed randomly in one of the 4 corners of an open plexiglass arena (60×40×35 cm) for 5 minutes. The total time spent in and the latency to entry the center of the arena (s) were recorded and scored using Anymaze.

2.7. Light-dark test.

The LD test was used to measure anxiety-like behavior (Bailey and Crawley, 2009). The animals were tested for 10 min in a light–dark rectangular box (60×40×35 cm) in which the aversive light compartment (40×40×35 cm) was illuminated by a 100 lux light. The dark

side (20×40×35 cm) had an opaque cover and ~ 0 lux of light. The two compartments were connected by an open doorway, which allowed the subjects to move freely between the two compartments. The test began by placing the animal in the dark compartment. The time spent in the light compartment (s), the latency to enter in the light chamber (s), and the total number of transitions, were measured using Anymaze.

2.8. Acoustic startle response and pre-pulse inhibition test.

The ASR and PPI tests were used to assess stress reactivity and sensorimotor gating functions. The tests were performed with a startle reflex measuring apparatus (SR-LAB; San Diego Instruments, San Diego, CA), as previously described (Toth et al., 2013). The system comprises a piezoelectric unit that transduces vibrations into signals when mice startle inside the plexiglas cylinder. First, mice are placed in a plexiglas cylinder with background noise (65 decibel [db]) for 5 min (acclimation phase). Then, mice were subjected to a total of 179 trials, for a total of 25 min, including: (a) startle trials (40 milliseconds [ms] with 80, 90, 100, 110 and 120 db acoustic pulses), (b) prepulse+startle trials: 20 ms with acoustic prepulses of 3 (68), 6 (71) and 12 (77) db above background noise followed, 100ms later, by a 40 ms 120-db startling pulse. Startle amplitude was measured every 1ms over a 65ms period beginning at the onset of the startle stimulus. Average startle amplitude over the sampling period was taken as the dependent variable. Percent PPI at each pre-pulse intensity was calculated as $100 - [(\text{startle response for prepulse} / \text{startle response for startle-alone trials}) \times 100]$.

2.9. Locomotor activity.

Locomotor activity was measured using the video tracking system Anymaze. Mice were placed in an empty open field (60×40×35 cm) for 20 minutes. The distance traveled (m) was recorded in 5 minutes intervals and used as an index of locomotor activity in a novel environment.

2.10. Body weight measurements.

Body weight (g) was measured through the course of the drug administration protocol and at PND63 when the behavioral assays began. The change in body weight was calculated as the difference between body weight at any given day and body weight at PND 28.

2.11. RNA Extraction, cDNA Synthesis and qPCR.

Brain tissue from PFC $n = 8$ female WT, $n = 6$ male WT, $n = 6$ female HR, $n = 5$ male HR mice was homogenized in TRIzol Reagent (#15596018, Thermo Fisher Scientific) and Zirconium beads (#Zr0B05-RNA, Next Advance) using the Bullet Blender homogenizer (BBX24B, Next Advance,). RNA was extracted on columns with the Direct-Zol RNA miniprep kit (#R2051, Zymo Research). To quantify *Reelin* expression levels, equal amounts of cDNA were synthesized using the Superscript VILO MasterMix (#11755-050, Thermo Fisher Scientific) and mixed with the qPCRBIO SyGreen Blue Mix (#17-507DB, PCR Biosystems) and 5 pmol of both forward (5'- GGACTAAGAATGCTTATTCC -3') and reverse (5'- GGAAGTAGAATTCATCCATCAG -3') *Reelin* primers. ACTB was amplified

as an internal control (5'- ATGGAGGGGAATACAGCCC -3') and reverse (5'- TTCTTTGCAGCTCCTTCGTT -3').

2.12. Plasma THC analysis.

For determination of plasma THC levels, male (n = 8) and female (n = 8) mice were used for each group. For the acute administration group, blood samples were collected 30 minutes after a single injection of THC at PND 48. For the chronic administration group, mice were injected with THC once a day for 21 consecutive days (PND 28–48) and blood samples were collected 24 hours after the last injection. Blood samples (~250 µL) were collected in tubes containing EDTA, via syringe needle insertion in the heart ventricles following exposure to carbon monoxide. Plasma THC concentration was quantified using fast liquid chromatography/mass spectrometry (LC/MS) adapted from (Lacroix and Saussereau, 2012; Irimia et al., 2015; Nguyen et al., 2018). 50 µL of plasma were mixed with 50 µL of deuterated internal standard (100 ng/mL CBD-d3 and THC-d3 in acetonitrile; Cerilliant), and cannabinoids were extracted from samples using 300 µL acetonitrile and 800 µL of chloroform, dried and then reconstituted in 100 µL of a methanol/water (2:1) mixture. Separation was performed on an Agilent LC1100 using a Poroshell 120 EC-C18 column (4.0µm, 2.1mm x 100mm) using isocratic elution with water and methanol, both with 0.2 % formic acid (250 µL/min; 81% MeOH). THC was quantified using an Agilent 6140 single quadrupole MSD using electrospray ionization and selected ion monitoring [THC (m/z=315.2) and THC-d3 (m/z=318.2)]. A linear scaling correction factor was used to quantify THC in one subject with 40 µL of plasma. Calibration curves were conducted daily for each assay at a concentration range of 0–200 ng/mL and observed correlation coefficients were 0.999.

2.13. Statistical analysis.

To examine how factors (treatment, genotype, sex) affected mice behavior or THC level, we used linear mixed models (LMM) in JMP pro v. 15.0 (SAS Institute, Inc). LMM allow modeling of both fixed and random effects (which subsume repeated measures) (Quinn GP, 2002; Zuur AF, 2016). We incorporated the categorical predictor variables of interest as fixed effects and included all possible interactions. We included mouse cohorts and individual subjects as random effects to account for possible non-independence of the data. We ensured assumptions of approximate normality and variance homogeneity were met by inspecting plots of residuals versus predicted values, and by inspecting quantile-quantile plots with 95% confidence limit curves. When residual plots indicated that it was appropriate for repeated measures, we used a covariance structure that allowed variances to differ across the levels of the repeated variable (Garrett M. Fitzmaurice, 2011). We analyzed the untransformed data in all but few traits where a fourth root transformation mitigated variance heterogeneity (e.g. latency to light, latency to center, Vmax of acoustic startle, body weight change). When significant interactions were found, planned post-hoc pairwise comparisons were performed to identify differences among specific genotype and treatment groups. We report *P*-values that remained significant after controlling for multiple comparisons by holding the 'false-discovery rate' to 0.05 using the Benjamini-Hochberg method (Hochberg, 1995). Outliers were detected using the Huber M-estimation method

(Huber, 1973) in JMP pro v. 15.0 and removed when appropriate ($n = 3$ in latency to light, $n = 2$ latency to center, $n = 2$ time spent sniffing, $n = 1$ sociability index, $n = 1$ in % PPI).

2.14. Factor analysis of behavioral assays.

Factor analysis is a data reduction method for understanding underlying relationships among variables (Bartholomew, 2008). We performed factor analysis using the variables from multiple behavioral assays, such as 6-DOT (DI, total exploration time in test trial [T3]), social interaction (time sniffing novel mouse, difference time sniffing novel mouse vs novel object), OF (time in center), LD (time in light), TS (mobility time). Factor analysis was computed in JMP 15.0 Pro and was conducted with varimax rotation with a factor-loading cutoff of 0.3 (Manly and Navarro Alberto; Stephens, 1996). The number of factors retained in our model was selected by inspecting the ‘elbow’ on the scree plot curve with factors retained if their eigenvalues were greater than 1 (Cattell, 1966). With these settings, a three-factor model was generated for our dataset. The loadings of the observed variables on the extracted factors are shown in Fig. 7. Factor scores for individual mice were extracted and used as variables for subsequent LMM analysis to identify their associations with treatment, sex, and genotype.

3. Results

3.1. Working memory was impaired after chronic adolescent exposure to THC, and in male HR mice

To study the long-term effects of chronic adolescent exposure to high doses of THC (10mg/kg), male and female mice were administered THC chronically during the adolescent period (PND 28–48) and were tested 2 weeks after the last injection of THC (Fig. 1A). Mice of both sexes showed similar level of THC in the plasma after acute or chronic injections (Fig. Suppl. 1A). To examine how Reelin signaling influences the behavioral effects of chronic adolescent exposure to THC, we used HR mice, which carry a null mutation in the *Reelin* gene (D’Arcangelo et al., 1995) and express ~40% reduced level of *Reelin* mRNA in brain tissues (Fig. 1B–C).

To assess the long-term effects of chronic adolescent exposure to THC on working memory in WT and HR mice, we used a modified novel object recognition test that uses 6 instead of 2 objects (6-DOT, Fig. 2A) (Sannino et al., 2012; Olivito et al., 2016). This test evaluates recognition memory under conditions of high loads of information processing, which is referred to as memory span and is considered a form of working memory. A discrimination index (DI) was calculated to determine the amount of time mice explored the novel object compared to the familiar objects. A preference for the novel object is considered as a sign that mice remember the familiar objects (Sannino et al., 2012). Higher rates of exploratory activity in response to the novel object is also interpreted as a sign of novelty seeking behavior (Flagel et al., 2014).

Chronic THC exposure during adolescence decreased the DI (working memory) of mice across all groups by ~31% compared to the vehicle-treated control group (Fig. 2B–C, treatment effect, $F_{1,86} = 16.4$, $P < 0.0001$). HR mice also showed decreased DI, but the

magnitude of the genotype effect varied among sexes (Fig. 2D, sex x genotype interaction, $F_{1,86} = 5.7$, $P = 0.019$). Compared to WT, female HR mice did not show impaired working memory, but male HR mice showed a decreased DI that was similar to the level of impairment caused by THC treatment in male WT mice (Fig. 2D, paired t test, $t = 2.9$, $P_{adj} = 0.019$).

This assay also revealed that HR mice displayed a ~25% increase in novelty-induced exploratory activity compared to WT controls, while treatment had no detectable effect (Fig. 2E–F, genotype effect, $F_{1,84} = 6.1$, $P = 0.015$; treatment effect $F_{1,81} = 2.3$, $P = 0.13$). This effect was triggered by the novel object, as it was revealed during the test phase while the total exploratory activity toward objects did not vary among groups in the initial familiarization phase (Fig. 2G, main effects and interactions P s > 0.3).

To further evaluate potential confounding effects of general locomotion or body weight having an impact on exploratory activity in this task, we analyzed changes in spontaneous locomotor activity in an empty arena or in body weight in WT and HR mice (Suppl. Fig. 1B–C). There was no effect of THC treatment, genotype, or sex on locomotor activity or body weight at the time of testing, suggesting that the deficits in working memory observed in WT and HR mice were not driven by changes in locomotion or body weight.

3.2. Social behaviors were impaired by chronic adolescent exposure to THC in male HR mice

To assess the long-term effects of chronic adolescent exposure to THC on social interaction behavior in WT and HR mice, we used the three-chamber social approach test in which a mouse is given the choice to spend time interacting with a novel mouse or a novel inanimate object, which are referred to as “exploration targets” in our analysis (Yang et al., 2011). We considered two aspects of social interaction: (1) time spent in the chamber with the novel mouse versus the novel object, and (2) time actively sniffing the novel mouse versus the novel object, which is a more specific measure of social interaction (Yang et al., 2011). Higher time spent in the chamber with or sniffing the novel mouse compared to the novel object was considered an index of “sociability”.

All mice spent longer time with novel mouse than the inanimate object (Fig. 3B–C, exploration target effect, $F_{1,119} = 49.5$, $P < 0.0001$), and sniffed the novel mouse more than the novel object (Fig. 3D–E, exploration target effect; $F_{1,113} = 83.3$, $P < 0.0001$), confirming that the behavioral assay can detect sociability across all groups. There were no significant effects of treatment, genotype and sex on time spent in the chamber (Fig. 3B, $P > 0.1$). In contrast, when analyzing time sniffing (Fig. 3D), the effect of chronic adolescent exposure to THC varied among the two exploration targets (Fig. 3F, treatment x exploration target interaction, $F_{1,113} = 4.1$, $P = 0.046$). Precisely, THC had no significant effect on time spent sniffing the novel inanimate object compared to vehicle (Fig. 3F, paired t test, $t = 0.06$, $P_{adj} = 1$); however, THC caused a ~20% decrease in time spent sniffing the novel mouse compared to vehicle across all groups (Fig. 3F, $t = 3.1$, $P_{adj} = 0.003$). The suppressive effect of THC on social interaction depended on the genotype (Fig. 3G, treatment x genotype interaction, $F_{1,108} = 6.1$, $P = 0.015$); a post-hoc analysis revealing that only HR mice treated with THC exhibited a significant reduction in overall time sniffing when compared to

vehicle treated HR mice (paired t test, $t = 3.2$, $P_{adj} = 0.006$). However, when a sociability index was calculated, we did not detect any effect of treatment or genotype (all effects and interaction P s > 0.1 , Suppl. Fig. 2).

3.3. Chronic adolescent exposure to THC increased anxiety-like behavior in female HR mice

To evaluate how impaired Reelin signaling influences anxiety-like behavior in mice chronically exposed to THC during adolescence, we compared WT and HR mice in the LD (Fig. 4A) and OF (Fig. 4F) tests. These behavioral assays examine anxiety-like responses by which animals avoid illuminated or open areas (Crawley and Goodwin, 1980; Bailey and Crawley, 2009). Decreased anxiety (increased disinhibition) is represented by mice spending more time in the light or open compartments, and by having a shorter latency to enter the brightly lit or center chambers.

In the LD test (Fig. 4B), THC increased the overall time spent in the light compartment (treatment effect, $F_{1,102} = 8.5$, $P = 0.0043$), but the strength of these effects varied among genotypes and sexes (Fig. 4C, treatment x genotype x sex interaction, $F_{1,97.5} = 4.4$, $P = 0.04$). This effect was mainly driven by female HR mice treated with THC, which spent 43% longer time in the light compared to the vehicle-treated group (Fig. 4C, paired t test, $t = 3.6$, $P_{adj} = 0.004$). A separate analysis revealed that the latency to enter the light chamber was also influenced by genotype and sex (Fig. 4D–E, sex x genotype interaction, $F_{1,47} = 5$, $P = 0.0197$). Female HR mice showed ~18% increase in latency compared to female WT, indicating an increased baseline anxiety-like responses (paired t test, $t = 2.5$, $P_{adj} = 0.03$). However, the latency to enter the light compartment was not significantly influenced by THC (treatment effect, $F_{1,97} = 0.02$, $P = 0.88$).

In the open field test (Fig. 4G), HR mice spent ~18% more time in the center (Fig. 4H, genotype effect, $F_{1,102} = 3.7$, $P = 0.057$ marginally significant), suggesting a reduced anxiety-like behavior. Time spent in center also varied among sexes and treatment groups (treatment x genotype x sex interaction, $F_{1,99} = 6.1$, $P = 0.016$); however, despite this significant overall variability, a post-hoc analysis did not identify significant differences between specific pairs of groups. Latency to enter the open area did not vary among groups (Fig. 4I, all $P > 0.2$).

3.4. Male HR mice showed enhanced ability to strive against stress following chronic adolescent exposure to THC

To test the effect of adolescent exposure to THC on reactivity to aversive conditions in WT and HR mice, we performed the TS test (Fig. 5A), an assay commonly used to screen anti-depressive drugs and to measure behavioral despair in mice (Can et al., 2012). An increased mobility has been associated with enhanced ability to strive against stress and is broadly related to active coping, impulsive and aggressive behaviors (Strekalova et al., 2004; Brockhurst et al., 2015).

We quantified total mobility time in a 6-minute tail suspension assay in WT and HR mice (Fig. 5B). THC treatment influenced total mobility time, but the effect varied among sexes and genotypes (Fig. 5C, treatment x sex x genotype interaction, $F_{1,158} = 12.8$, $P = 0.0005$).

While THC did not significantly change the mobility time of either male or female WT mice compared to the vehicle controls (paired t test, $t = 1.9$, $P_{adj} = 0.07$; $t = 0.5$, $P_{adj} = 0.6$, respectively), it induced a 19% increase in mobility time for male HR mice (Fig. 5C, paired t test, $t = 2.6$, $P_{adj} = 0.04$).

3.5. Reelin deficiency led to enhanced startle responses and pre-pulse inhibition

To evaluate psychotic-like behaviors in WT and HR mice following chronic adolescent exposure to THC, we quantified the ASR and the % PPI (Fig. 6A). The ASR is a reflexive reaction to a sudden acoustic stimulus (Swerdlow et al., 2001; Geyer et al., 2002). The % PPI is used as a measure of sensorimotor gating, which occurs when a weak pre-pulse stimulus suppresses the response to the subsequent startling stimulus. PPI is impaired in schizophrenia patients and serves as an animal model of schizophrenia (Cadenhead et al., 1993).

We first examined the differences in ASR of WT and HR mice at various acoustic intensities (65, 80, 90, 100, 110, 120 db, Fig. 6B). THC did not influence ASR (treatment effect, $F_{1,451} = 0.7$, $P = 0.4$). Compared to WT, HR mice exhibited an overall ~3% higher startle response (genotype effect, $F_{1,451} = 7.2$, $P = 0.008$), which varied by the acoustic intensity of the pulse (Fig. 6C, genotype x db interaction, $F_{5,408} = 2.8$, $P = 0.016$). The increased startle tended to be stronger at 120db (Fig. 6B, paired t test, $t = 2$, $P_{adj} = 0.08$ marginally significant).

When examining the % PPI at three pre-pulse intensities (Fig. 6D, 68, 71 and 77 db), HR mice showed an overall ~13% increase in % PPI compared to WT mice (Fig. 6E, genotype effect, $F_{1,482} = 8.6$, $P = 0.004$). Adolescent exposure to THC also had an effect on % PPI that varied among sexes (Fig. 6F, treatment x sex interaction, $F_{1,482} = 8$, $P = 0.005$). Female mice treated with THC tended to show ~12 % reduction in % PPI compared to vehicle-treated mice (Fig. 6F, paired t test, $t = 2$, $P_{adj} = 0.05$).

3.6. THC and Reelin deficiency influenced behavioral domains that were associated with cognition and disinhibition

We performed a factor analysis to determine whether the multiple behavioral responses We performed a factor analysis to determine whether the multiple behavioral responses influenced by THC and *Reelin* deficiency.

The factor analysis identified three unique behavioral domains (factors), which explained 57% of total variance (Fig. 7A). The first factor explained 23.8% of the total variance and contained outcomes (positive loadings) from the social interaction and working memory tests (e.g., time sniffing novel mouse, discrimination index). As factor 1 largely reflected measures of cognitive functions, we named it “*cognition*”. The second factor explained 15.7% of the total variance and was named “*stress reactivity*” because the main outcome that contributed to this cluster was the mobility time of the TS test. The third factor (17% of the total variance) was named “*disinhibition*” as it contained behaviors that reflected increased exploratory or disinhibited behavior (e.g., time in the center of the open field, novel object exploration).

To determine whether these behavioral domains differed between treatments, genotypes, or sexes, we analyzed the rotator factor scores for each subject. We found that the behavioral domains were significantly influenced by treatment or genotype (Fig. 7B, treatment x behavioral domains interaction, $F_{2,142} = 5.7$, $P = 0.004$; Fig. 7C, genotype x behavioral domains interaction, $F_{2,142} = 3.4$, $P = 0.035$; respectively). Post-hoc analysis revealed that *cognition* was significantly decreased by THC (Fig. 7B, paired t test, $t = 3.1$, $P_{adj} = 0.006$), and *disinhibition* was significantly increased by *Reelin* deficiency (Fig. 7C, paired t test, $t = 2.8$, $P_{adj} = 0.014$). Neither treatment, genotype, nor sex significantly influenced the third behavioral domain, *stress reactivity*.

Overall, this exploratory data analysis is consistent with the results of the separate behavioral tests by indicating that adolescent exposure to THC detrimentally affected cognitive functions, and it revealed that reduced expression of *Reelin* led to disinhibitory behaviors, which were intensified by THC treatment in a sex-specific manner.

Discussion

The present study revealed for the first time that reduced levels of *Reelin* influences behavioral abnormalities caused by heavy consumption of THC during adolescence, in a sex-dependent manner. Here we discuss the potential implications and limitation of our findings.

Influence of THC and Reelin deficiency on cognitive functions

In line with previous reports in rodent models and clinical studies in humans, we found that chronic adolescent exposure to THC impaired working memory (Renard et al., 2014; Rubino and Parolaro, 2016; Hurd et al., 2019). Additionally, HR male mice showed lower working memory in comparison to the WT littermates and to a similar extent as did chronic adolescent exposure to THC. Although it is the first time that HR mice are tested in a task that measures memory span, such as the 6-DOT, this finding is consistent with studies demonstrating the crucial role of *Reelin* in mechanisms underlying learning and memory (Weeber et al., 2002; Rogers et al., 2013; Iafrati et al., 2014; Telese et al., 2015). Overall, these findings show that our experimental model reproduced previously published results; however, the lack of a genotype x treatment interaction in the 6-DOT suggests that the underlying mechanisms may be independent.

We found that social interaction was impaired by chronic adolescent exposure to THC with the strongest effects in HR mice. These observations are in line with evidence in human and preclinical studies showing social deficits induced by THC (Long et al., 2010). Only few studies have examined social behaviors of the HR mice and did not report prominent social deficits (Podhorna and Didriksen, 2004; Macri et al., 2010; Michetti et al., 2014). Thus, our study revealed for the first time that social interaction is reduced by THC in mice with *Reelin* deficiency. It will be important to examine if these behavioral changes are also associated with cellular and molecular changes in areas implicated in social behavior, including PFC, amygdala and striatum (Ko, 2017).

Consistently, the factor analysis revealed that variables from both working memory and social interaction tests contributed to a behavioral domain, named ‘*cognition*’, which was negatively influenced by adolescent exposure to THC. The negative effects of THC on cognitive behaviors are known to be mediated by CB1 receptors in rodents (Lichtman and Martin, 1996; Nava et al., 2001; Niyuhire et al., 2007; Wise et al., 2009; Puighermanal et al., 2013). Therefore, it is reasonable to infer that the behavioral effects induced by THC in our study are dependent on CB1 receptors; however, future research is necessary to confirm this.

Reelin deficiency is associated with disinhibitory phenotypes

We speculate that multiple behavioral responses exhibited by HR mice may reflect general behavioral disinhibition. In rodents, behavioral disinhibition is measured as a function of increased exploratory activity towards unfamiliar objects or environments, and has been associated with compulsive drug taking (Davis et al., 2008; Flagel et al., 2010; Belin et al., 2011). In humans, behavioral disinhibition reflects personality traits that encompasses impulsivity, risk-taking and novelty seeking phenotypes; these behavioral patterns are more pronounced in adolescence and have been linked to addiction susceptibility (Young et al., 2009). Here, we found that, compared to WT, both male and female HR mice explored the novel objects for longer time in the 6-DOT. Additionally, female HR mice treated with THC showed signs of reduced inhibitory control as they spent more time in the center chamber of the OF test or in the brightly lit compartment of the LD test. Consistently, variables from these tests contributed to the same behavioral domain (‘*disinhibition*’) revealed by the factor analysis, which was positively influenced by *Reelin* deficiency. These observations are also in line with a previous study showing decreased anxiety of HR mice in the elevated plus maze and increased motor impulsivity (Ognibene et al., 2007).

Collectively, these results suggest that Reelin deficiency leads to loss of inhibitory control, which may underlie behavioral traits linked to addiction vulnerability. Further supporting this hypothesis, in a recent genome-wide association study (GWAS), a variant of the *Reelin* gene was associated with higher likelihood to consume alcohol ($P = 4 \times 10^{-9}$, variant and risk allele rs756747-T) (Karlsson Linner et al., 2019). It will be important to experimentally test whether HR mice show increased drug taking behavior when exposed to different drugs of abuse.

Reelin deficiency leads to abnormal responses to aversive conditions

We observed abnormal behavior in HR mice in response to aversive conditions. First, male HR mice treated with THC showed prolonged mobility time in the TS test compared to vehicle group. This behavior has been linked to proactive coping in response to a threat (Strekalova et al., 2004; Brockhurst et al., 2015). Second, HR mice exhibited enhanced startle reactivity to acoustic stimuli compared to WT mice, which was not influenced by THC treatment. In a previous study, the startling response of HR mice did not change following a single acoustic pulse of 105db; however, in our study, we used five startling pulses from 80db to 120db that likely increased the sensitivity of the task. Elevated ASR has been observed in mental conditions associated with impaired emotional reactivity, such as posttraumatic stress disorders (Orr et al., 1995) and obsessive-compulsive disorders (Kumari et al., 2001). These observations led us speculate that reduced expression of *Reelin* may lead

to altered emotional reactivity. This hypothesis deserves further examination of the HR mice using tasks specifically designed to assess impulsivity and aggression.

Reelin as a susceptibility factors for psychiatric disorders

The array of behavioral phenotypes exhibited by HR mice in our study, encompassing memory impairments, social deficits, poor inhibitory control, and altered stress responses, is reminiscent of the behaviors characterizing numerous psychiatric conditions, ranging from autism spectrum disorders (ASD) to schizophrenia and substance use disorders.

Consistently, a role of Reelin in the development of these disorders is supported by several lines of evidence. In particular, whole exome sequencing studies identified *de novo* mutations in the *Reelin* gene in individuals with ASD (Neale et al., 2012; Iossifov et al., 2014; Wang et al., 2014). Further support for Reelin involvement in psychiatric disorders is provided by the observation of reduced expression of *Reelin* transcript or protein in postmortem brains of individuals affected by ASD (Fatemi et al., 2001), and schizophrenia (Guidotti et al., 2000; Ruzicka et al., 2007; Habl et al., 2012). These observations suggest that altered Reelin signaling may be a vulnerability factor for psychiatric disorders and that the HR mice represent a valuable animal model for translational research. However, we did not observe any PPI deficits in HR mice. In contrast, a previous study showed that a single *in vivo* injection of Reelin protein increases % PPI (Rogers et al., 2013). It is possible that, in our study, the elevated startle response observed in HR mice may act as a confounding factor for accurately assessing the effect of *Reelin* deficiency. It is also possible that impaired Reelin signaling contributes mainly to the cognitive, social and emotional deficits associated with schizophrenia, which are referred to as negative symptoms, as opposed to positive symptoms that include psychotic-like behaviors (Kirkpatrick et al., 2006).

Notably, THC reduced % PPI in female WT mice, suggesting that a history of drug exposure leads to psychotic-like behaviors. Conflicting results have been reported concerning the effects of adolescent chronic exposure to THC on psychotic-like behaviors in rodents (Rodriguez et al., 2017; Todd et al., 2017; Ibarra-Lecue et al., 2018). Differences in THC doses, mice genetic background and specific experimental conditions among different research groups may explain some of these conflicts.

Sex differences in behavioral responses associated with Reelin haploinsufficiency

Our study revealed numerous sex differences in the behavioral abnormalities associated with *Reelin* deficiency and/or adolescent exposure to THC. Because we did not observe sex differences in the plasma levels of THC following acute or chronic THC administration, we conclude that other factors are responsible for driving sex differences in the observed behavior. Male HR mice were more sensitive to working memory impairments, as well as to emotional reactivity in the TS test. In contrast, female mice, showed reduced anxiety-like behaviors (HR) or reduced % PPI (WT) in response to THC. Consistently, previous finding in rodents showed that chronic adolescent exposure to cannabinoids is associated with numerous behavioral sex differences; however, the knowledge of the underlying mechanisms remains limited (Craft et al., 2013). Given the critical role of Reelin brain development, it is possible that Reelin influences the development of the endocannabinoid system in a sex-specific manner, which could lead to sex differences in the behavioral effects of THC.

Limitations of the experimental model

Our study focused on the adolescent exposure to THC due to its relevance to human studies showing that cannabis is the most widely used illicit drug among adolescents. We explicitly note that the observed effects of adolescent exposure may also occur following chronic adult exposure, which remains to be investigated.

Our study used a single dosage of THC (10mg/kg) and did not examine overall dose response curves, which would permit better understanding of potential sex and genotype differences. However, this dosage can elevate plasma levels of THC in rodents similar to those detected in humans using cannabis; hence, it should adequately model heavy chronic THC exposure, which have been linked to detrimental health effects. Specifically, we show that 30 minutes after a single injection of THC (10 mg/kg) in mice the plasma concentrations of THC were similar to those reported in humans after smoking marijuana or vaping ethanolized (pure) THC (>100ng/ml) (Abrams et al, 2007; Zuurman et al, 2008, Huestis et al, 1992, Heustis et al, 2007). Moreover, we show that 24 hours after the last daily injection of chronic (21 days) THC treatment in mice the plasma levels of THC were similar to those detected in cannabis frequent users (> 6ng/ml) (Desrosiers et al., 2014; Lee et al., 2015).

In conclusion, our study indicates that Reelin deficiency contributes to behavioral alterations induced by THC exposure and suggests that elucidating Reelin signaling will improve our understanding of neurobiological mechanisms underlying behavioral traits relevant to the development of psychiatric conditions.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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- Decreased anxiety-like behavior was observed in female HR mice treated with THC
- Enhanced stress reactivity was observed in male HR mice

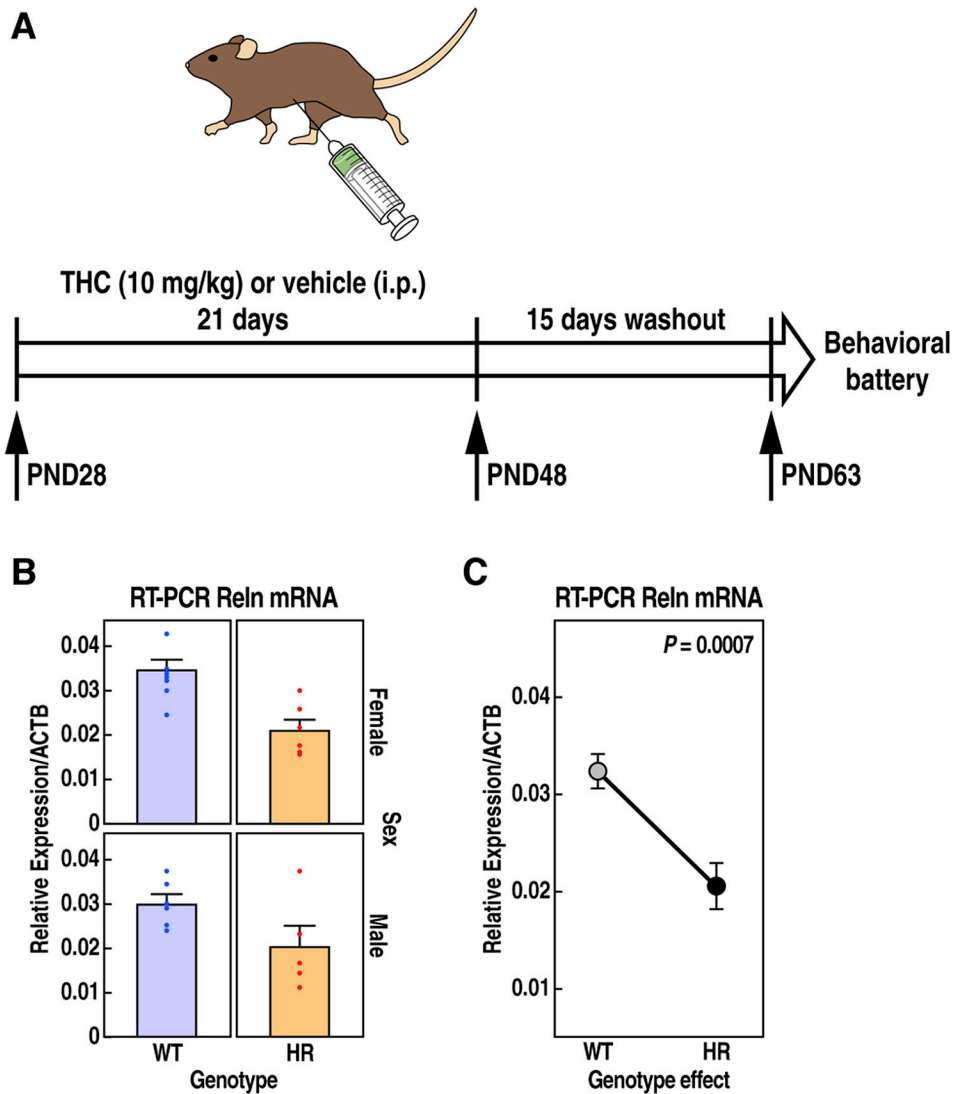


Figure 1: Experimental model.

(A) Schematic diagram illustrates the daily i.p. administration of THC (10 mg/Kg) during adolescence (PND28–48), and the day when behavioral assessment began (PND63) after 15 days of washout. (B) Relative expression of Reelin mRNA transcript levels measured by RT-PCR and normalized to housekeeping gene ACTB in total RNA isolated from mouse brain tissues in WT and HR mice. (C) Reelin expression is reduced in HR mice compared to WT littermates (main genotype effect, $P = 0.0007$, LMM).

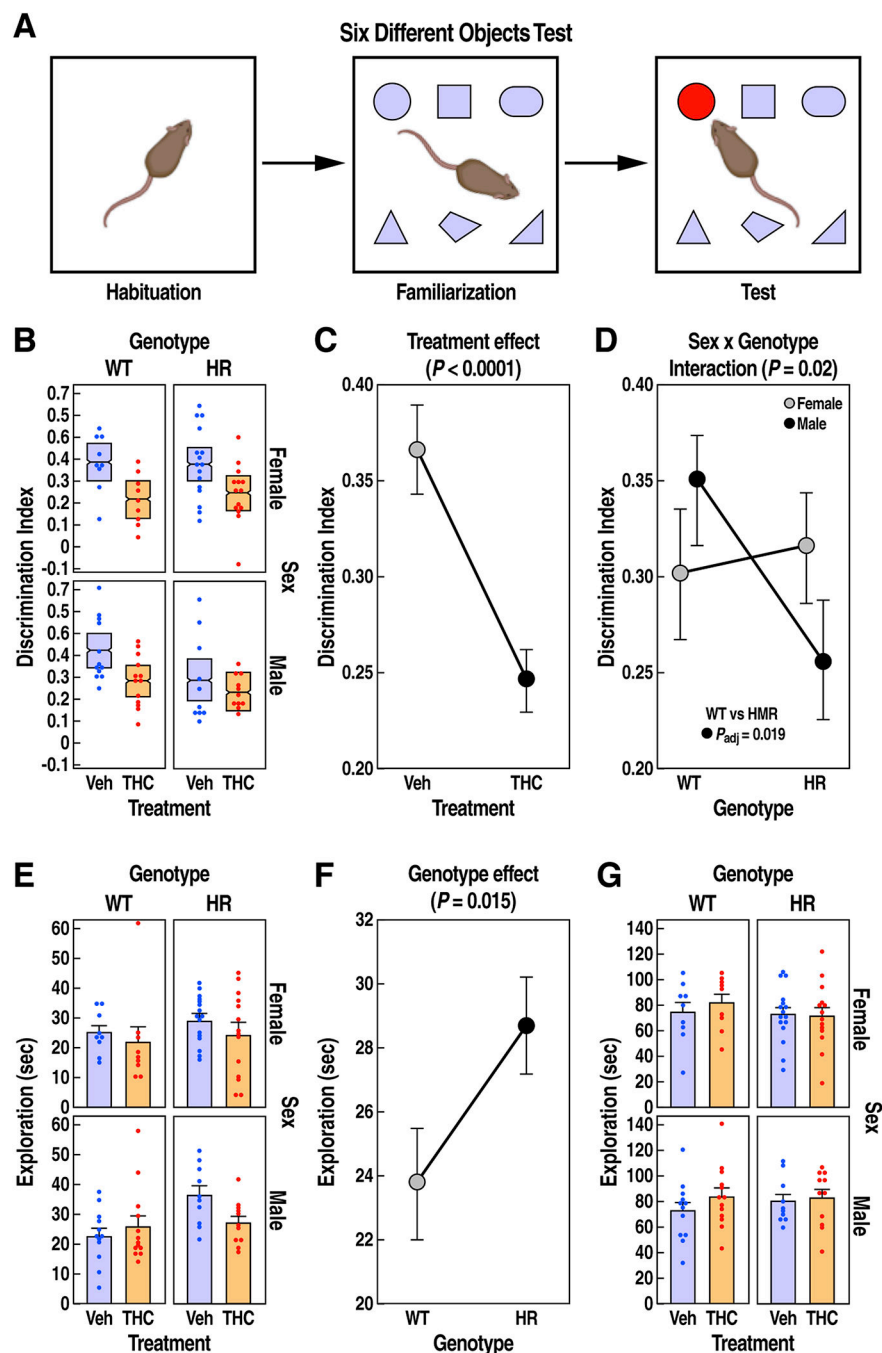


Figure 2: Six-different objects test in WT and HR mice after chronic adolescent exposure to THC.

(A) 6-DOT with habituation, familiarization, and test trials. (B) DI is shown as mean \pm 95% confidence intervals. (C) THC reduced the DI (mean \pm SEM) across all groups (main treatment effect, $P < 0.0001$, LMM). (D) Male HR mice showed a reduced DI (mean \pm SEM) compared to male WT mice (sex x genotype interaction, $P = 0.02$, LMM). (E) Exploration time (s) in test trial is expressed as mean \pm SEM. (F) HR mice spent longer time (mean \pm SEM) exploring the objects in the test trial (main genotype effect, $P = 0.015$,

LMM). (**G**) Exploration time (mean \pm SEM) in familiarization trial did not change across groups. FDR adjusted $P < 0.05$ for post-hoc pairwise t tests are reported.

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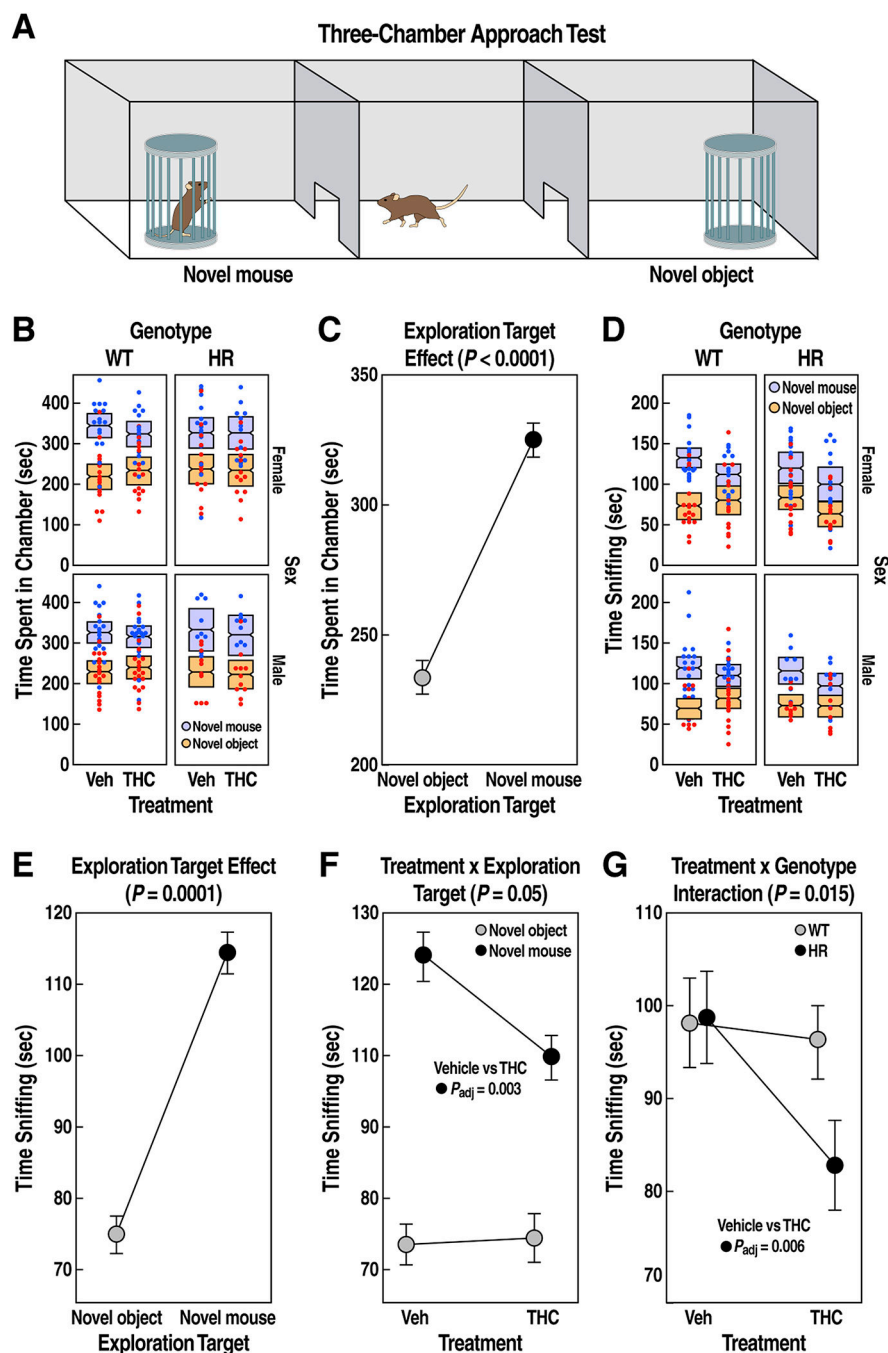


Figure 3: Social interaction behavior in WT and HR after chronic adolescent exposure to THC. (A) Three-chamber interaction test diagram. Time (sec) spent in chamber (B) or sniffing (D) is expressed as mean \pm 95% confidence intervals. (C) Mice spent more time (mean \pm SEM) in the chamber or (D) more time sniffing (mean \pm SEM) the novel mouse compared to the novel object (main exploration target effect, $P < 0.0001$, LMM). (E-F) THC reduced social investigation (post-hoc pairwise t test $P_{\text{adj}} = 0.003$; treatment x exploration target interaction, $P = 0.05$, LMM). (G) HR exposed to THC during adolescence showed reduced social

interaction behavior (post-hoc pairwise t test $P_{\text{adj}} = 0.006$; treatment x genotype interaction, $P = 0.015$, LMM). FDR adjusted $P < 0.05$ for post-hoc pairwise t tests are reported.

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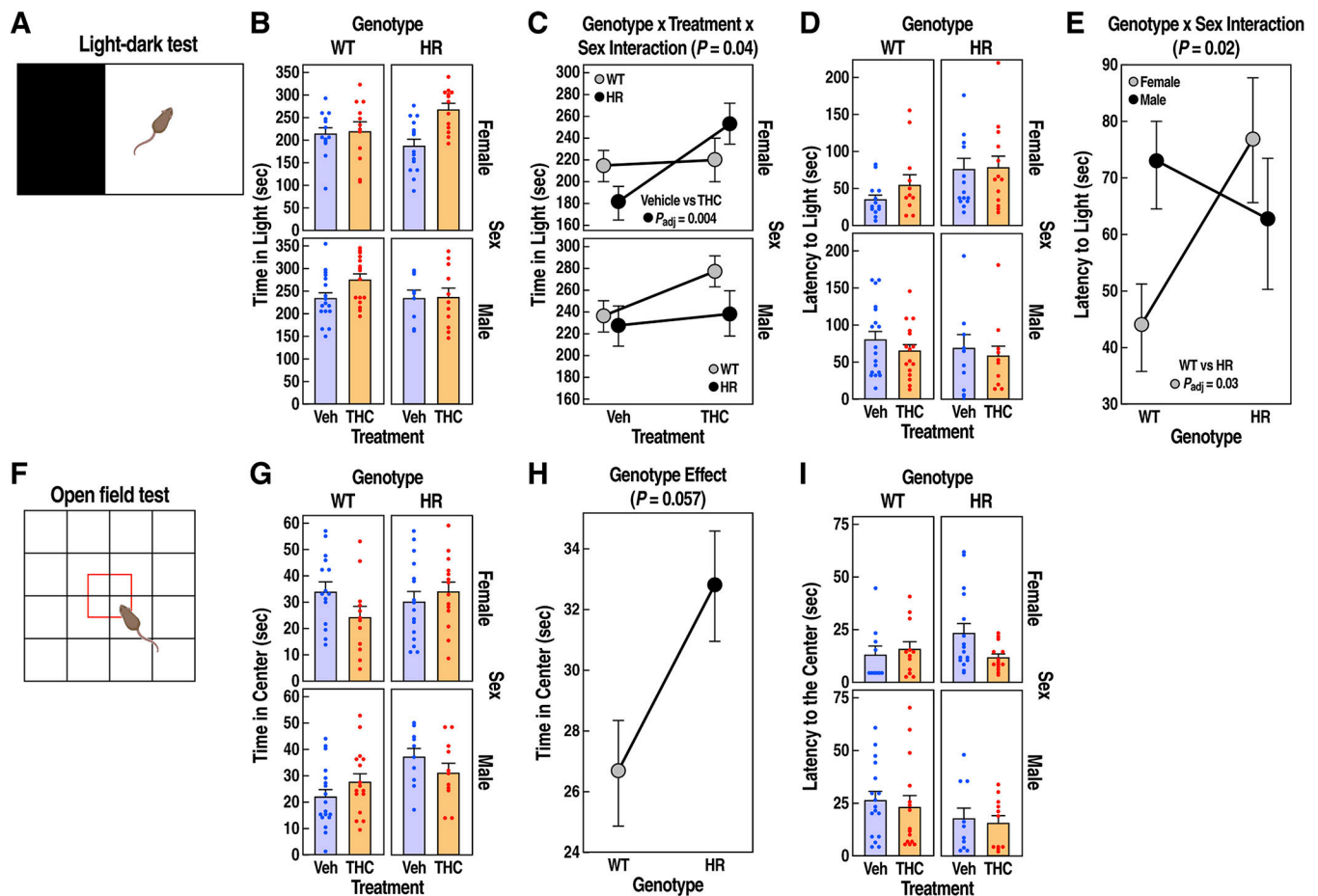


Figure 4: Anxiety-like responses in WT and HR following chronic adolescent exposure to THC. (A) Light-dark test schematic. (B) Time (sec) spent in the light compartment is expressed as mean \pm SEM. (C) Female HR treated with THC during adolescence showed reduced anxiety (treatment x genotype x sex interaction, $P = 0.04$, LMM). (D) Latency (sec) to enter the illuminated chamber is expressed as mean \pm SEM. (E) Female HR mice showed increased latency (sec) to enter the light compared to WT controls (genotype x sex interaction, $P = 0.02$, LMM). (F) Open field test schematic. (G) Time (sec) spent in the center of the arena is expressed as mean \pm SEM. (H) HR mice showed reduced anxiety (main genotype effect, $P = 0.057$, LMM). (I) Latency (sec) to enter the center compartment is expressed as mean \pm SEM and did not change across groups. FDR adjusted $P < 0.05$ for post-hoc pairwise t tests are reported.

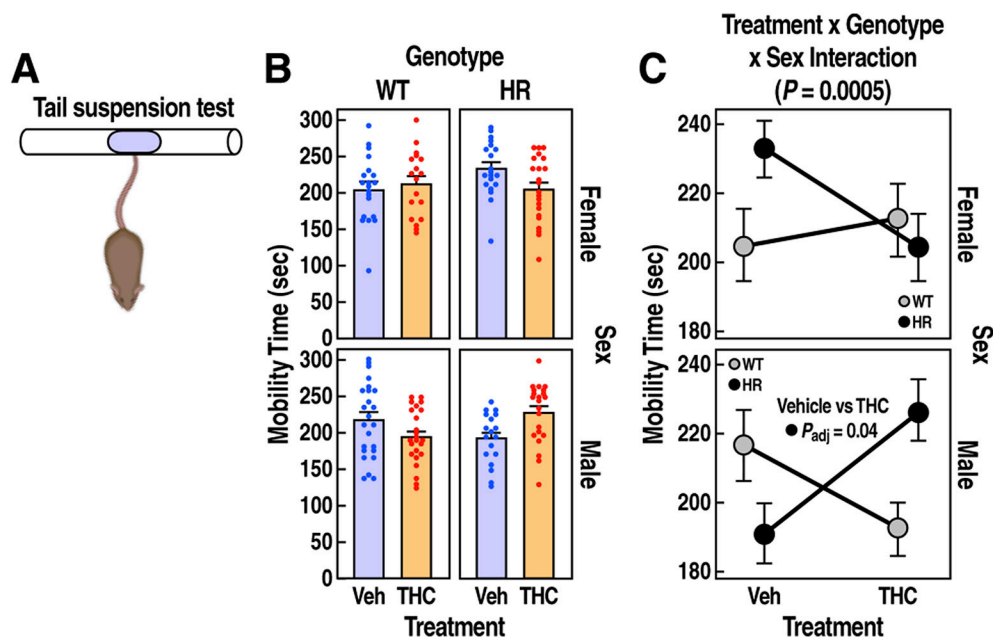


Figure 5: Reactivity to aversive conditions in WT and HR mice following chronic adolescent exposure to THC.

(A) Tail suspension test diagram. (B) Mobility time (sec) is expressed as mean \pm SEM. (C) Male HR mice treated with THC during adolescence showed increased mobility (treatment x genotype x sex interaction, $P = 0.0005$, LMM). FDR adjusted $P < 0.05$ for post-hoc pairwise t tests are reported.

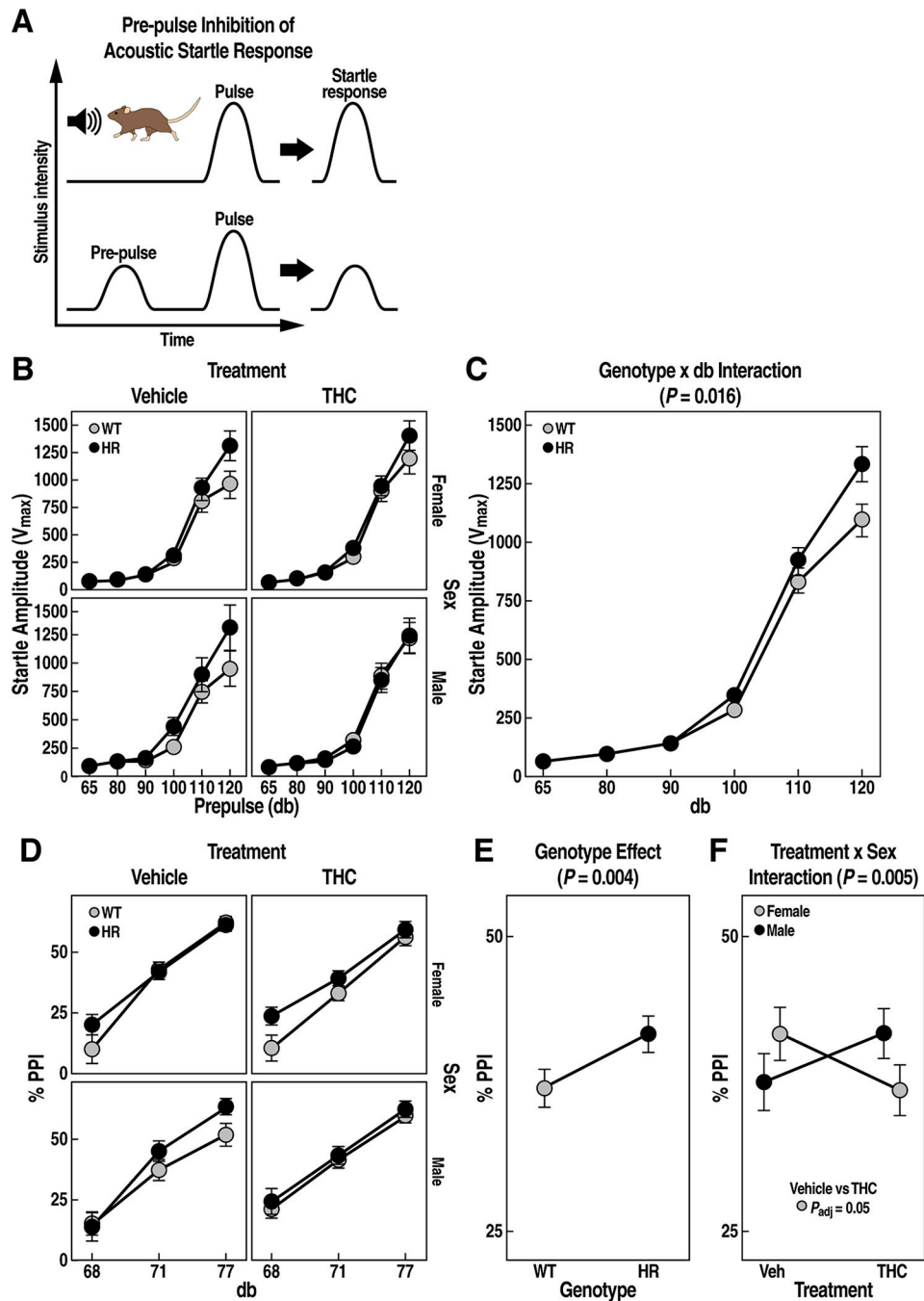


Figure 6: Acoustic startle reflex of pre-pulse inhibition in WT and HR mice following chronic adolescent exposure to THC.

(A) Pre-pulse inhibition of acoustic startle test diagram. (B) Startle amplitude is expressed as $V_{max} \pm SEM$. (C) HR mice show increased startle response (genotype x db interaction, $P = 0.016$, LMM). (D) Pre-pulse inhibition is expressed as $\% PPI \pm SEM$. (E) HR mice exhibited higher $\% PPI$ compared to WT controls (main genotype effect, $P = 0.004$, LMM). (F) Female treated with THC during adolescence showed reduced $\% PPI$ compared to

vehicle (treatment x sex interaction, $P = 0.005$, LMM). FDR adjusted $P < 0.05$ for post-hoc pairwise t tests are reported.

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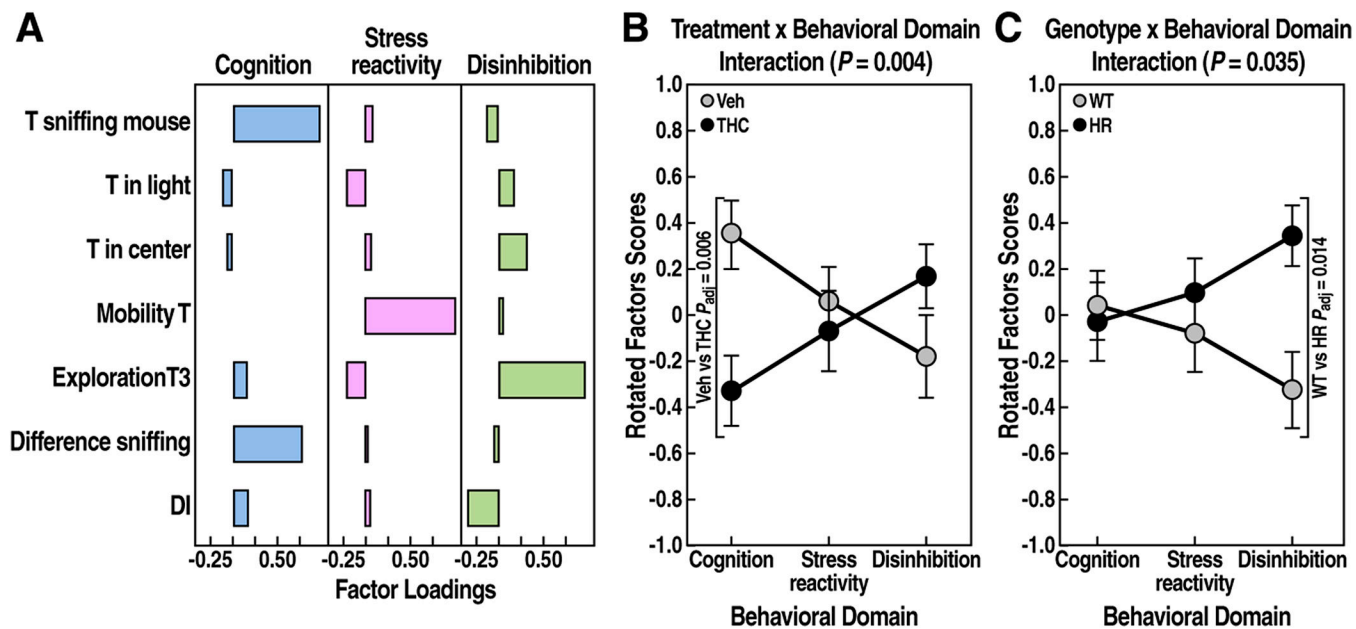


Figure 7: Factor analysis of behavioral tests.

(A) Factor loadings for each named behavioral domain are reported for several behavioral variables. LMM analysis for rotated factor scores revealed a (B) treatment x behavioral domain interaction ($P = 0.004$), and a (C) genotype x behavioral domain interaction ($P = 0.035$). FDR adjusted $P < 0.05$ for post-hoc pairwise t tests are reported.

Table 1.

Number of mice used for behavioral analysis*

BEHAVIORAL ASSAYS	MALE WT		FEMALE WT		MALE HET		FEMALE HET	
	VEH	THC	VEH	THC	VEH	THC	VEH	THC
Locomotor	19	20	19	18	17	21	19	23
Tail suspension	24	24	20	18	18	22	20	22
Open field	17	15	10	12	10	11	16	14
Light-dark	18	16	13	12	9	11	15	13
Social	19	19	17	15	10	11	16	14
6-DOT	12	13	9	9	10	11	16	14
PPI	23	24	24	23	20	22	25	26
ASR	23	24	25	23	20	22	25	26

*The number of outliers for specific variables and removed from analysis of are reported in the method section

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