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Chronic antipsychotic treatment exerts limited effects on the mania-like behavior of dopamine transporter knockdown mice

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Abstract

Background: Bipolar disorder has been linked to dopamine transporter (DAT) polymorphisms and reduced DAT levels in postmortem brains.

Aims: The purpose of this study was to examine the effects of D_2 -family receptor antagonists on DAT dysfunction-mediated mania behavior in mice.

Methods: DAT knockdown mice received either D_2 -family receptor antagonist risperidone or asenapine and mania-related behaviors were assessed in the behavioral pattern monitor.

Results: Chronic risperidone did not reverse mania-like behavior in DAT knockdown mice. Chronic asenapine reduced mania behavior but this effect was more pronounced in wildtype littermates than in DAT knockdown mice.

Conclusion: Taken together, these findings suggest that while acute antipsychotic treatment may be beneficial in management of bipolar mania, it may not be necessary or effective to continue chronic antipsychotic treatment after an acute mania episode has been reversed. Further, it may be useful to explore whether DAT polymorphisms can predict chronic antipsychotic nonresponse in bipolar disorder mania.

Keywords

bipolar disorder; mania; dopamine transporter; behavioral pattern monitor; antipsychotic; mice

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Introduction

Bipolar disorder (BD) is a frequently disabling, life-threatening mental illness that occurs in >2.5% of the global population. BD is characterized by striking and persistent mood fluctuations ranging from mania to depression, driving clinically significant impairment (Grande et al., 2016). Current treatments include antidepressants and mood stabilizers, with more recent use of antipsychotics (Sachs, 2003). Today, a majority of patients with BD are prescribed antipsychotics (Ventimiglia et al., 2009). Not all treatments are effective in all patients however, and the neurobiological specificity of actions of these drugs remains to be elucidated. While mood stabilizers can vary in mechanism, the common mechanism for antipsychotics is dopamine D_2 receptor blockade (Stroup et al., 2003). Determining whether such blockade remediates specific behaviors associated with mania would prove beneficial to tailor patient-targeted treatments.

Neurobiological and genetic studies have implicated dopamine dysfunction in the etiology of BD. Impaired dopamine homeostatic mechanisms may result in extreme shifts to and from hyperdopaminergia and hypodopaminergia possibly driving mania and depressive states (Ashok et al., 2017). Altered dopaminergic states could arise from altered homeostatic clearance mechanisms, such as reduced levels of the dopamine transporter (DAT), which removes dopamine from the synaptic cleft. Reduced DAT levels were reported in unmedicated people with BD euthymia (Anand et al., 2011), potentially arising from genetic polymorphism-induced reduction in DAT expression (Greenwood et al., 2006; Horschitz et al., 2005). Targeted reduction of DAT expression results in seasonal sensitivity consistent with BD (Young et al., 2018), while DAT knockdown (KD) mice recreate many behavioral aspects of BD mania (Young et al., 2011; Greenwood et al., 2006; Pinsonneault et al., 2011; Zhuang et al., 2001) with a high fidelity of reproducibility such as in the cross-species behavioral pattern monitor (BPM) (Kwiatkowski et al., 2017). Specifically in the BPM, both BD mania patients and DAT KD mice exhibit hyperactivity, increased exploration, and altered locomotor patterns including straighter patterns of movement (reduced spatial d) (Perry et al., 2009), behaviors that are consistent over time (Minassian et al., 2011). Further, acute DAT blockade with GBR12909 consistently induces such behaviors, and the hyperactivity of both models can be reduced with mood stabilizers such as lithium or valproate (van Enkhuizen et al., 2013). Determining whether selective dopamine D₂ receptor blockade could more completely remediate such mania symptoms from reduced DAT function is important.

Hence, we tested the effects of chronic treatment with two atypical antipsychotics in the mouse behavioral pattern monitor (BPM) using DAT KD mice. We predicted that blockade of dopamine D_2 -family receptors would affect the BD mania model mice more strongly than controls, attenuating mania-like exploratory behavior.

Materials and methods

Animals

All animals were group housed (max. 4 per cage) and maintained in a temperaturecontrolled vivarium ($21 \pm 1^{\circ}$ C) on a reversed day-night cycle (lights on at 1900 hours and

off at 0700 hours). All mice had ad libitum access to food and water. Testing was performed during the dark phase of the day-night cycle between 0800 and 1800 hours. All procedures were approved by the UCSD Institutional Animal Care and Use Committee. The UCSD animal facility meets all federal and state requirements for animal care. Male and female adult DAT KD and WT littermates on C57BL/6J background were used to study the effects of chronic risperidone and chronic asenapine in the BPM.

Drug Treatment

Drug solutions were prepared daily and chronically administered for 28 days, and also 60 minutes prior to testing on the last day. Risperidone (Sigma-Aldrich, St. Louis, MO) was dissolved in water at concentrations of 0.03 mg/kg/day and 0.3 mg/kg/day. Risperidone doses were determined based upon previous research (Duncan et al., 2006). Asenapine maleate powder was obtained from Lundbeck and dissolved in saline at 0.03 and 0.1 mg/kg/day, injected at a volume of 5 ml/kg and administered via osmotic minipumps (Alzet, type 2004). Asenapine doses were determined based on previous studies (Ene et al., 2015). See Table 1 for the experimental design.

Behavioral Pattern Monitor

The primary measures investigated in the current study were transitions and center entries representing locomotor activity, holepoking and rearing representing exploratory behavior, and spatial d representing locomotor patterns. Spatial d measures geometric patterns of locomotor activity – i.e., straight line movements versus more circumscribed paths of movement. A spatial d value closer to 1 reflects a straighter path, and values closer to 2 indicating highly circumscribed small-scale movements (Kwiatkowski et al., 2017).

Plasma concentrations of asenapine and risperidone were determined from trunk blood taken after treatment and decapitation using ultra performance liquid chromatography (UPLC) coupled to tandem mass spectrometry (MS/MS). Plasma samples were precipitated with acetonitrile and DMSO (80:20 v/v) containing internal standard. Following centrifugation, 10 ul was injected onto the chromatographic system (Acquity UPLC system; Waters, Milford, MA). MS detection was performed on a Sciex-API 400 mass spectrometer (AB

Sciex. Foster City, CA). The lower limit of quantification was 5 ng/ml and 1 ng/ml for asenapine and risperidone.

Data analysis

Data were analyzed using a two-way or three-way analysis of variance (ANOVA) with drug, genotype, and sex as between subject factors. Significant main effects and interactions were followed up with one-way ANOVAs as well as Tukey *post hoc* analyses. The data were analyzed for 60 min testing periods using the BMDP statistical software (Statistical Solutions Inc., USA). The alpha was set at 0.05.

RESULTS

Chronic risperidone effects on exploration in DAT KD and WT mice

Risperidone on activity—Male and female mice were used in this experiment. A main effect of genotype was observed on all measures of activity, including counts $[F_{(1,79)}=107.56, p<0.0001]$, transitions $[F_{(1,79)}=98.38, p<0.0001]$, center entries $[F_{(1,79)}=51.66, p<0.0001]$, distance travelled $[F_{(1,79)}=112.81, p<0.0001]$, with KD mice more active than WT littermates. There was an effect of sex on activity as measured by counts $[F_{(1,79)}=4.01, p<0.05]$ and distance travelled $[F_{(1,79)}=4.05, p<0.05]$ and a trend toward an effect of sex on transitions $[F_{(1,79)}=3.39, p<0.1]$, with males exhibiting higher levels than females. No sex x genotype interactions were observed.

Risperidone treatment exerted a main effect on center entries $[F_{(2,79)}=3.89, p<0.05]$, with a trend effect on counts $[F_{(2,79)}=2.51, p<0.1]$ and distance travelled $[F_{(2,79)}=2.63, p<0.1]$. No genotype x drug interactions were observed, but there was a significant sex x drug x genotype interaction on transitions $[F_{(2,79)}=3.25, p<0.05]$, as well as a trend toward a significant sex x drug x genotype interaction on center entries $[F_{(2,79)}=5.59, p<0.01]$ and distance travelled $[F_{(2,79)}=3.01, p<0.1]$. No other interactions were observed. Post-hoc analyses revealed that for center entries DAT KD mice were not different compared to WT vehicle-treated animals at the highest risperidone dose, suggesting possible attenuation of the DAT KD effect at this dose (see Figure 1, Table 3).

Risperidone on exploration—We examined the effects of sex, genotype, and risperidone on inspective and diversive exploration. There were significant effects of genotype on all measures of inspective and diversive exploration – holepoking $[F_{(1,79)}=14.57, p<0.0005]$, repeated holepoking $[F_{(1,79)}=13.56, p<0.0005]$, varied holepoking $[F_{(1,79)}=13.86, p<0.0005]$, rearing $[F_{(1,79)}=30.46, p<0.0001]$, and center duration $[F_{(1,79)}=5.31, p<0.05]$. There was a trend toward an effect of risperidone on center duration $[F_{(1,79)}=2.60, p<0.1]$. There was a significant effect of sex on rearing $[F_{(1,79)}=11.09, p<0.005]$, with males exhibiting more rearing. There was a significant interaction effect of sex x genotype on varied poking $[F_{(1,79)}=4.72, p<0.05]$, There were no other significant interaction.

Post-hoc analyses revealed a trend toward an increase in holepoking in DAT KD vehicle and DAT KD 0.03 mg/kg risperidone-treated animals compared to WT 0.3 mg/kg risperidone-treated animals, and a significant increase in DAT KD 0.3 mg/kg risperidone-treated

animals compared to WT 0.3 mg/kg risperidone (p<0.05), suggesting risperidone had a larger effect on reducing holepoking in WT animals compared to DAT KD animals. For varied holepoking, there was a trend toward an effect of DAT KD 0.03 mg/kg risperidone compared to WT 0.3 mg/kg risperidone and a significant increase in DAT KD 0.3 mg/kg risperidone compared to WT 0.3 mg/kg risperidone, suggesting high dose risperidone may have opposite effects on varied holepoking in DAT KD mice compared to WT mice, and only decreases varied holepoking in WT mice at the doses studied. For repeated holepoking, DAT KD vehicle, DAT KD 0.03 mg/kg risperidone and DAT KD 0.3 mg/kg risperidone treated mice showed a trend toward an increase compared to WT 0.3 mg/kg risperidone, again demonstrating differential effects of risperidone in WT and DAT KD mice. Post-hoc analyses revealed an increase in rearing in DAT KD mice compared to WT littermates and this effect was most pronounced when comparing DAT KD 0.3 mg/kg risperidone mice to WT 0.3 mg/kg risperidone treated mice (p < 0.01), suggesting this antipsychotic has a general sedating effect but does not specifically ameliorate the underlying issue of DAT dysfunction (see Figure 1). Center duration was increased in DAT KD 0.03 mg/kg risperidone-treated animals compared to WT 0.3 mg/kg risperidone-treated animals (p < 0.05), suggesting less aversion to risk in risperidone-treated DAT KD mice compared to risperidone-treated WT mice.

Risperidone on locomotor patterns—There were significant effects of genotype on all locomotor pattern variables – spatial d $[F_{(1,79)}=19.53, p<0.0001]$, entropy h $[F_{(1,79)}=93.97, p<0.0001]$, spatial CV $[F_{(1,79)}=4.08, p<0.05]$, temporal CV $[F_{(1,79)}=11.80, p<0.001]$. There were no effects of sex on locomotor patterns. There were no significant effects of risperidone on locomotor pattern, nor were there any significant interaction effects on locomotor pattern. Post-hoc analyses revealed a significant decrease in spatial d in KD DAT vehicle treated mice compared to WT vehicle treated mice (p<0.05) and WT 0.3 mg/kg risperidone (p<0.01). A priori hypothesis testing revealed a trend toward a decrease in KD DAT 0.3 mg/kg risperidone mice compared to WT vehicle-treated and KD DAT 0.03 mg/kg risperidone-treated mice compared to WT 0.3 mg/kg risperidone-treated mice. Post-hoc analyses revealed a mice compared to WT vehicle-treated and KD DAT 0.03 mg/kg risperidone-treated mice compared to WT 0.3 mg/kg risperidone-treated mice. Post-hoc analyses revealed a mice compared to WT vehicle-treated mice. Post-hoc analyses revealed an increase in entropy *h* in all DAT KD groups compared to all WT groups (p<0.01). There were no significant post-hoc results for spatial CV. There was an increase in temporal CV in WT 0.3 mg/kg risperidone mice compared to DAT KD vehicle (trend), 0.03 mg/kg risperidone (p<0.01) and 0.3 mg/kg risperidone (trend) treated animals.

Chronic asenapine effects on exploration in DAT KD and WT mice

Asenapine on activity—Only male mice were studied in this experiment. There was a significant effect of genotype on activity as reflected by increased counts $[F_{(1,60)}=134.7, p<0.0001]$, transitions $[F_{(1,60)}=77.2, p<0.0001]$ and distance travelled $[F_{(1,60)}=24.64, p<0.0001]$. There was a main effect of asenapine on counts $[F_{(2,60)}=11.8, p<0.0001]$ and transitions $[F_{(2,60)}=4.9, p<0.05]$ while an interaction between genotype and asenapine was only observed for counts $[F_{(2,60)}=6.1, p<0.005,$ see Figure 2], but not transitions. Post-hoc analyses revealed that WT mice receiving the highest dose of asenapine (1 mg/kg/day) exhibited fewer counts compared with vehicle-treated WT mice (p<0.05).

Asenapine on exploration—DAT KD mice in this experiment did not exhibit increased holepoking, repeated holepoking or varied holepoking but reared more $[F_{(1,60)}=9.7, p<0.005]$ than WT mice. Asenapine tended to affect holepoking $[F_{(2,60)}=2.5, p<0.1]$ and did affect rearing $[F_{(2,60)}=11.1, p<0.0005]$. Genotype and asenapine tended to interact in holepoking $[F_{(2,60)}=3.0, p<0.1]$, while significant interactions were observed on rearing $[F_{(2,60)}=4.7, p<0.05]$ (see Figure 2). Post-hoc analyses revealed that WT mice receiving the highest dose of asenapine (1 mg/kg/day) tended to exhibit fewer holepokes (p<0.1) and exhibited fewer rears (p<0.05) compared with WT vehicle-treated mice.

Asenapine on locomotor patterns—DAT KD mice exhibited more linear movements as measured by lower spatial $d[F_{(1,60)}=9.0, p<0.005]$ and more erratic patterns of movement as measured by higher entropy h $[F_{(1,60)}=7.6, p<0.0001]$ compared with WT mice. Asenapine did not affect spatial d (F<1.8, ns) but did lower entropy h, increasing the predictability of movement $[F_{(2,60)}=7.6, p<0.005]$ (see Figure 2, Table 4). Genotype and asenapine did not interact in any measure of locomotor patterns (F<2, ns).

Plasma concentrations of risperidone and asenapine in DAT KD and WT mice

Plasma levels of risperidone (N = 25) and asenapine (N = 15) were quantified in DAT KD and WT mice. Plasma levels of risperidone and asenapine were not significantly different in DAT KD mice compared to WT mice (p=0.17 and 0.42, respectively; see Table 5). In BD mania when used in concert with other drug treatment, the recommended dose of asenapine is 5 mg twice a day (Marazziti et al., 2016), and plasma asenapine levels in the current study were much higher than what is seen in humans following 5 mg asenapine treatment (below 6 ng/mL) (Gerrits et al., 2012). The recommended dose of risperidone in bipolar mania is 2-3 mg once daily (Sajatovic et al., 2006). In humans, 2 mg risperidone was associated with plasma levels around 3 ng/mL at 8 weeks, which is higher than what we observed in DAT KD mice but lower than what was observed in WT mice (Cardoni, 1995).

Discussion

DAT KD mice exhibited hyperexploratory patterns consistent with patients with BD mania. Overall, chronic antipsychotic treatment with risperidone or asenapine was ineffective in fully normalizing mania-like behaviors at the doses tested. This finding contrasts with previous findings that acute antipsychotic treatment attenuated hyperactivity, hyperexploration, and lower spatial d of DAT KD mice (Milienne-Petiot et al., 2017) as well as findings that chronic valproate attenuated the hyperactivity of DAT KD mice (van Enkhuizen et al., 2013) and chronic lithium attenuated the hyperactivity and hyperexploratory behavior of GBR12909-treated mice (van Enkhuizen et al., 2015). Our findings are consistent with acute antipsychotic treatment responsiveness in acute mania episodes but suggest chronic antipsychotic treatment may not be efficacious as a long-term prophylactic treatment for mania.

Previous work suggests short-term but not long-term antipsychotic treatment reverses prepulse inhibition deficits and inhibits amphetamine and tail-pinch induced hyperlocomotion in C57BL6 mice (Amato et al., 2018). The authors found the decline in antipsychotic efficacy was correlated to extracellular dopamine levels rather than reductions

in D₂ receptor occupancy by antipsychotics. Moreover, antipsychotic treatment failure is associated with a progressively linearized relationship between TH and DAT, which was present after long- but not short-term treatment (Amato et al., 2018). Dopamine receptor densities following chronic treatment also have been observed (Silvestri et al., 2000) and may play a role in changes in efficacy of antipsychotics over time in managing BD mania. Thus, while acute antipsychotic treatment is useful for BD mania, and chronic for maintenance in BD euthymia, chronic treatment-induced dysregulation of dopamine clearance may limit efficacy for mania treatment.

An alternate explanation could be that the compounds administered were biologically inert. This possibility remains unlikely because plasma levels were in a range comparable to those seen in humans at dosages used to manage BD mania. In addition, several main effects of treatments were observed in these studies.

Although asenapine failed to normalize mania-like behavior in DAT KD mice at the doses investigated, asenapine impacted WT behavior. Chronic asenapine treatment reduced activity, exploration, and reduced straight line movements and predictability of movement in WT mice specifically at the highest dose, but this effect was attenuated in DAT KD mice. Therefore, in terms of exploratory behavior, DAT KD mice were less sensitive than WT mice to chronic asenapine treatment. This observation indicates that DAT genotype may predict non-response to antipsychotics. Risperidone partially reduced center entries, increased rearing, and had a small effect in decreasing overall counts. Plasma risperidone and asenapine levels were quantified and although reduced antipsychotic levels were observed in DAT KD vs. WT mice, this effect was not statistically significant. This observation likely rules out differences in drug metabolism between genotypes, although larger sample sizes would be preferable. Nevertheless, higher risperidone and asenapine doses may be warranted to correct mania-relevant behaviors DAT KD mice.

In both experiments DAT KD mice exhibited mania-like behaviors, although these behaviors were more widespread in experiment 1 compared to 2, likely as a result of background strain differences between C57BL/6 and s129 strains (Kwiatkowski et al, 2018). Female DAT KD mice exhibit BD mania-relevant behaviors. Although the estrogen cycle in female mice may affect behavior, these effects are not large enough to justify excluding female mice from mouse behavioral research (Kokras and Dalla, 2014). BD type II affects women more frequently than men, while BD type I affects women and men at equal rates (Diflorio and Jones, 2010). There were no sex x genotype interactions for any variable except for varied poking, in which female DAT KD mice exhibited the most varied poking. This observation is in contrast to previous work in our group showing no sex effects on any measure of the BPM. This inconsistency may be related to use of a different background strain in the previous work (Young et al., 2010), although findings across strains have been consistent for the majority of measures as shown in a meta-analysis (Kwiatkowski et al 2018).

The results of this study indicate that it may be useful to examine whether DAT polymorphisms can predict antipsychotic nonresponse in bipolar disorder. Based on previous work, it is possible that DAT polymorphisms may instead predict response to valproate. Valproate increases DAT mRNA and protein via increased H3K9 and Nurr1 enrichment in

the DAT promoter (Green et al., 2017). In addition, valproate has been shown to reduce hyperactivity and improve restricted locomotor patterns in DAT KD mice (van Enkhuizen et al., 2013). Interestingly, increased dopamine in DAT KD mice has been shown to be associated with a lengthening of circadian rhythm, while valproic acid reverses this pattern (Landgraf et al., 2016). Chronic valproate treatment also has been shown to attenuate GBR 12909-induced hyperactivity, without affecting other behavioral parameters such as specific exploration or sequential organization (van Enkhuizen et al. 2013a).

Dysfunctional dopamine homeostasis has been linked to many diverse psychiatric illnesses and is not limited to BD. Moreover, dopamine dysfunction can be influenced by a variety of factors, including genetic factors such as DAT polymorphisms (Herborg et al., 2018) (Pinsonneault et al., 2011) as well as environmental factors such as childhood adverse events or trauma (Pruessner et al., 2004) (van Winkel et al., 2013). Notably, PTSD comorbidity with BD is associated with poorer clinical outcome (Quarantini et al., 2010). Understanding the contributors to dopamine dysregulation within subjects may be a first step to identifying the correct treatment, which may require multifaceted treatment approaches such as gene-specific pharmaceuticals, nutraceuticals, cognitive behavioral therapy, vagus nerve stimulation, and/or diet. These interventions would be determined at an individual level to best address comorbid conditions and readouts of specific dopamine dysfunction, which could potentially be exacerbated by multiple variables, not simply a single gene.

The results of this study support DAT polymorphisms as a target for pharmacogenetic studies of predictors of drug response in BD. Research on pharmacogenetics of lithium response in bipolar disorder is ongoing (Papiol et al., 2018; Hou et al., 2016; McCarthy et al., 2010), while investigation of pharmacogenetics of antipsychotic response in bipolar disorder has largely been extrapolated from research in subjects with schizophrenia. Importantly, these data support the idea that antipsychotic drugs may provide clinical benefit acutely, but may not provide additional benefit when administered chronically in BD. It is therefore prudent to conduct a benefit-risk assessment of chronic antipsychotic treatment for bipolar disorder mania and to determine if it may be beneficial to cease antipsychotic treatment following the acute mania episode.

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Figure 1. Risperidone effects on BPM performance in DAT KD and wildtype mice.

There were significant effects of genotype on all measures. There was a significant effect of risperidone on center entries. There were no significant genotype x risperidone interactions. Abbreviations: WT – wildtype littermates; DAT KD – dopamine transporter knockdown; Veh – vehicle; 0.03 Risp – 0.03 mg/kg risperidone; 0.3 Risp – 0.3 mg/kg risperidone. Data presented as Mean +/– SEM. Statistically significant differences compared to wildtype vehicle controls are denoted with * or ** (p < 0.05 and p < 0.01, respectively). Statistically significant main effects of gene are denoted with #.



Figure 2. Asenapine effects on BPM performance in DAT KD and wildtype mice.

There were significant effects of genotype on all measures except for center entries. There was a significant effect of asenapine on counts, rearing, and entropy h. There were significant genotype x asenapine interactions on counts and rearing, and this was related to a decrease in counts and rearing in wildtype mice. Abbreviations: WT – wildtype littermates; DAT KD – dopamine transporter knockdown; Veh – vehicle; 0.03 Asen – 0.03 mg/kg asenapine; 0.1 Asen – 0.1 mg/kg asenapine). Data presented as Mean +/– SEM. Statistically significant differences compared to wildtype vehicle controls are denoted with * or ** (p < 0.05 and p < 0.01, respectively). Statistically significant main effects of gene are denoted with #.

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Experimental design for experiments 1 and 2, DAT=Dopamine Transporter, KD=knockdown

Experiment 1 – Chronic ris DAT KD mice on a C57BL/	peridone in 5-6 6J background (month old (male/female)
Drug Treatment	<u>Wildtype</u>	DAT KD
Vehicle	6/10	8/5
Risperidone 0.03 mg/kg	6/10	10/6
Risperidone 0.3 mg/kg	6/10	8/6
Experiment 2 – Chronic as DAT KD mice	mapine in 3 mor	nth old male
Drug Treatment	Wildtype	DAT KD
Vehicle	6	13
Asenapine 0.03 mg/kg	6	12
Asenapine 0.1 mg/kg	10	13

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Measurements and their associated factors in the mouse Behavioral Pattern Monitor.

Factor	Measure	Description
	Counts	The cumulative number of distinct behaviors to occur during testing
A straight and an	Transitions	The number of movements from one pre-defined area to another
ACUVILY INTERSULTES	Center Entries	The number of entries into the area defined as the center
	Distance travelled	Total distance travelled while exploring the chamber measured in cm
	Hole pokes	The total number of investigatory holepokes across all 11 holes
	Repeated Poking	The number of times a hole was repeatedly poked before moving to another hole
Specific Exploration	Varied Poking	The number of times the mouse holepoked into different holes
	Rear	The number of times the mouse reared into the air
	Center Duration	The amount of time (s) spent in the area designated as the center of the chamber
	Spatial d	Spatial scaling exponent d measures the hierarchical and geometric organization of behavior (Paulus, Callaway, & Geyer, 1993)
Locomotor Patterns	Entropy h	Quantifies the sequential aspects of sequences of behavior, specifically measuring the degree to which behavior is observed along a continuum between complete order and disorder (Paulus, Geyer, & Braff, 1996)
	Spatial CV	Measures the consistency by which the animal moves from one region to another (Geyer, Russo, & Masten, 1986)
	Temporal CV	Measures the consistency of the dwell time in each of the regions (Geyer et al., 1986)

Table 3.

Risperidone, DAT KD genotype and sex effects on BPM activity.

Significant findings in bold, trend toward significance in italics.

	Measure	Sex df (1,79)	Genotype df (1,79)	Drug df (2,79)	Sex x Genotype df (1,79)	Sex x Drug df (2,79)	Genotype x Drug df (2,79)	Sex x Drugs x Genotype df (2,79)
	Counts	F=4.01, p=0.05	F=107.56, p=0.00	F=2.51,p=0.09	F=0.01,p=0.92	F=0.50, p=0.61	F=1.14,p=0.33	F=2.23,p=0.11
A activite.	Transitions	F=3.39, p=0.07	F=98.38, p=0.00	F=2.23,p=0.11	F=0.02,p=0.89	F=0.75, p=0.48	F=0.51, p=0.60	F=3.25,p=0.04
ACUVILY	Center Entries	F=1.57,p=0.21	F=51.66, p=0.00	F=3.89,p=0.02	F=0.00,p=0.98	F=2.03, p=0.14	F=0.35,p=0.71	F=5.59,p=0.01
	Distance travelled	F=4.05, p=0.05	F=112.81,p=0.00	F=2.63,p=0.08	F=0.12,p=0.73	F=0.93, p=0.40	F=0.67, p=0.51	F=3.01,p=0.05
	Hole pokes	F=0.65, p=0.42	F=14.57,p=0.00	F=0.34,p=0.71	F=3.07,p=0.08	F=0.77,p=0.47	F=0.81, p=0.45	F=1.39,p=0.25
Inspective Exploration	Repeated Poking	F=0.54, p=0.46	F=13.56,p=0.00	F=0.36,p=0.70	F=2.57,p=0.11	F=0.80, p=0.46	F=0.68, p=0.51	F=1.61,p=0.21
	Varied Poking	F=0.99,p=0.32	F=13.86,p=0.00	F=0.14, p=0.87	F=4.72,p=0.03	F=0.42,p=0.66	F=1.54, p=0.22	F=0.51, p=0.60
Dirouciro Funlouotion	Rearing	F=11.09, p=0.00	F=30.46, p=0.00	F=0.12,p=0.88	F=0.07,p=0.79	F=0.85,p=0.43	F=1.03,p=0.36	F=0.27,p=0.76
	Center Duration	F=0.51, p=0.48	F=5.31,p=0.02	F=2.60,p=0.08	F=0.15, p=0.70	F=0.81, p=0.45	F=0.21,p=0.81	F=1.46,p=0.24
	Spatial d	F=0.20, p=0.66	F=19.53,p=0.00	F=0.51, p=0.60	F=0.48,p=0.49	F=0.54, p=0.58	F=1.42,p=0.25	F=0.27,p=0.76
I accurator Dattorne	Entropy h	F=0.05, p=0.82	F=93.97,p=0.00	F=1.42,p=0.25	F=0.15, p=0.70	F=1.11,p=0.33	F=0.65, p=0.53	F=2.57,p=0.08
	Spatial CV	F=0.00, p=0.95	F=4.08,p=0.05	F=2.23,p=0.11	F=1.98,p=0.16	F=0.37,p=0.69	F=0.09,p=0.91	F=1.03,p=0.36
	Temporal CV	F=0.02,p=0.90	F=11.80,p=0.00	F=0.84,p=0.44	F=1.46,p=0.23	F=0.43,p=0.65	F=0.55,p=0.58	F=0.62,p=0.54

Table 4.

Effects of asenapine and DAT KD genotype on BPM activity.

Activity	Measure	Cellotype	Asellapille	
Activity		df (1,60)	df (2,60)	df (2,60)
Activity	Counts	F=134.65,p=0.00	F=11.78,p=0.00	F=6.07, p=0.00
ACUVIU	Transitions	F=77.20,p=0.00	F=4.90,p=0.01	F=1.47,p=0.24
)	Center Entries	F=0.70,p=0.41	F=1.04,p=0.36	F=1.34, p=0.27
Di	istance travelled	F=24.64,p=0.00	F=2.48, p=0.09	F=0.10,p=0.91
	Hole pokes	F=1.48,p=0.23	F=2.45, p=0.10	F=3.02, p=0.06
Inspective Exploration R	Repeated Poking	F=1.28,p=0.26	F=1.95, p=0.15	F=2.61, p=0.08
	Varied Poking	F=1.60,p=0.21	F=4.13,p=0.02	F=3.42,p=0.04
Dirondiro E malouotion	Rearing	F=9.66, p=0.00	F=11.14,p=0.00	F=4.67, p=0.01
	Center Duration	F=54.09, p=0.00	F=1.47,p=0.24	F=0.43, p=0.66
	Spatial d	F=9.03, p=0.00	F=1.77,p=0.18	F=0.17, p=0.84
Locomoton Dottome	Entropy h	F=75.11,p=0.00	F=7.59, p=0.00	F=1.99, p=0.15
	Spatial CV	F=102.57, p=0.00	F=8.08, p=0.00	F=2.10,p=0.13
-	Temporal CV	F=0.28, p=0.60	F=4.31,p=0.02	F=2.64, p=0.08

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Risperidone and asenapine blood levels were quantified in WT and DAT KD mice. Means +/- Standard error are shown (ng/mL).

	Risperidone	Asenapine
WТ	4.9 + - 0.6 (n = 23)	26 +/- 13 (n =7)
DAT KD	2.0 +/- 0.7 (n = 2)	15.5 +/- 2.9 (n = 8)