scientific reports

OPEN



Male and female behavioral variability and morphine response in C57BL/6J, DBA/2J, and their BXD progeny following chronic stress exposure

Carole Morel^{1,2,6}, Lyonna F. Parise^{2,3,6}, Yentl Y. Van der Zee^{2,3,6}, Orna Issler^{2,3}, Min Cai¹, Caleb J. Browne^{2,3}, Anthony Blando¹, Katherine B. LeClair^{2,3}, Antonio V. Aubry^{2,3}, Sherod Haynes¹, Robert W. Williams⁴, Megan K. Mulligan⁴, Scott J. Russo^{2,3 \boxtimes}, Eric J. Nestler^{2,3 \boxtimes} & Ming-Hu Han^{1,2,5 \boxtimes}

Drug addiction is a multifactorial syndrome in which genetic predispositions and exposure to environmental stressors constitute major risk factors for the early onset, escalation, and relapse of addictive behaviors. While it is well known that stress plays a key role in drug addiction, the genetic factors that make certain individuals particularly sensitive to stress and, thereby, more vulnerable to becoming addicted are unknown. In an effort to test a complex set of gene x environment interactions—specifically *gene x chronic stress*—here we leveraged a systems genetics resource: BXD recombinant inbred mice (BXD5, BXD8, BXD14, BXD22, BXD29, and BXD32) and their parental mouse lines, C57BL/6J and DBA/2J. Utilizing the chronic social defeat stress (CSDS) and chronic variable stress (CVS) paradigms, we first showed sexual dimorphism in social and exploratory behaviors between the mouse strains. Further, we observed an interaction between genetic background and vulnerability to prolonged exposure to non-social stressors. Finally, we found that DBA/2J and C57BL/6J mice preexposed to stress displayed differences in morphine sensitivity. Our results support the hypothesis that genetic variation influences chronic stress-induced behavioral outcomes such as social and approachavoidance behaviors, reward responses, as well as morphine sensitivity, and is likely to modulate the development of drug addiction.

Drug addiction is a partly heritable, polygenic disorder determined by a complex interaction between multiple genes and the environment¹. Familial, twin, and adoption studies have consistently reported that genetic factors contribute to aspects of addiction primarily through interactions with environmental factors and exposure^{2,3}. Converging lines of evidence from both clinical and preclinical investigations support the view that environmental stress is a major risk factor for developing drug addiction^{4–6}. Indeed, clinical studies strongly suggest that cumulative or prolonged stress is a reliable predictor of drug addiction⁵, and preclinical studies in rodents have confirmed that chronic exposure to stress increases initiation of drug use, induces more robust drug-induced conditioned place preference (CPP), and escalates drug self-administration^{5,7,8}.

Although genetic studies have shown addiction to be roughly 50% heritable⁹⁻¹³, opioid addiction is accelerating in the US, and we are faced with a worldwide problem that accounts for tremendous morbidity and mortality, thereby posing collateral damage to social and political systems^{14–17}. Still, the DNA variants that contribute to the

¹Department of Pharmacological Sciences, Icahn School of Medicine at Mount Sinai, New York, NY, USA. ²Friedman Brain Institute, and Center for Affective Neuroscience, Icahn School of Medicine at Mount Sinai, New York, NY, USA. ³Nash Family Department of Neuroscience, Icahn School of Medicine at Mount Sinai, New York, NY, USA. ⁴Department of Genetics, Genomics and Informatics, University of Tennessee Health Science Center, Memphis, Tennessee, USA. ⁵Brain Cognition and Brain Disease Institute, Shenzhen Institute of Advanced Technology (SIAT), and Department of Mental Health and Public Health, Faculty of Life and Health Sciences, Shenzhen University of Advanced Technology (SUAT), Chinese Academy of Sciences, Shenzhen, Guangdong, People's Republic of China. ⁶These authors contributed equally: Carole Morel, Lyonna F. Parise and Yentl Van der Zee. ^{\Box}email: scott.russo@mssm.edu; eric.nestler@mssm.edu; ming-hu.han@mssm.edu behavioral relationship between stress and addiction remain elusive. To this end, the development of expanded families of fully isogenic replicable cohorts^{18–20}, in particular, the extended BXD family of recombinant inbred mouse strains generated from a cross between C57BL/6J females and DBA/2J males, have enabled large research efforts in systems genetics and have been used widely to investigate genetic factors underlying complex heritable phenotypes observed in metabolic^{21–23} and psychiatric disorders^{24–26}. Each of the ~120 extant BXD strains are essentially immortal allowing for reliable and reproducible genetic investigations. The BXD family provides a unique system to map the complex set of gene x environment (GxE) interactions, such as those between genes and stress that are implicated in the vulnerability to drug addiction^{26–31}.

In the present study, we first exposed male and female C57BL/6J, DBA/2J, BXD5, BXD8, BXD14, BXD22, BXD29 and BXD32 mice to the chronic social defeat stress paradigm (CSDS)^{32,33}. We identified social interaction and exploratory behavior variabilities amongst BXD strains and sexes after CSDS. To investigate the interaction between genetics and vulnerability to non-social stressors, we also exposed C57BL/6J, DBA/2J, BXD8, BXD22, and BXD29 male and female mice to the chronic variable stress paradigm (CVS)³⁴. We established that novelty-suppressed feeding, sucrose preference and exploratory behaviors after CVS differ among BXD progeny and by sex. Finally, female DBA/2J and C57BL/6J mice pre-exposed to CSDS displayed differences in morphine sensitivity. Characterization of the genetic, social, and environmental factors mediating addiction risk will fundamentally improve our understanding of individual variations in response to stress and drug abuse and provide highly useful information for developing new treatments for psychiatric disorders.

Results

C57BL/6J, DBA/2J, and BXD mice have distinct behavioral performances following chronic social stress

Drug addiction, including morphine-related behaviors, has been linked to several factors such as sex, age, social environment, and stress experience prior to drug exposure⁵. Similar to humans, mice exposed to chronic social stress exhibit a higher propensity to develop addictive-like behaviors. In our study, we first aimed to determine the effect of genetics, including sex divergence, on chronic social stress-induced behavioral outcomes. Here we used CSDS, a well-establish mouse model for social stress-induced behavioral alterations^{32,35}.

We exposed adult male and female mice from the BXD founders, C57BL/6J and DBA/2J, and the BXD5, BXD8, BXD14, BXD22, BXD29, and BXD32 strains to CSDS (Fig. 1a). We assessed their social interaction behavior determined by the social interaction ratio (see detailed information in the methods section). We first tested the null hypothesis that behavioral performance was independent of the mice's genetic profile, including sex chromosomes as genetic divergence. We observed a large range of social interaction ratios across mouse lines and sex (Fig. 1b; One-Way ANOVA, $F_{(15,223)}$ =3.805, p<0.001; see Supplementary Table S1 for details). We further observed an interaction between sex and genetic background (Two-Way ANOVA, $F_{(1,225)}$ =5.573, p<0.001). Following social interaction, we tested exploratory behaviors in an open-field test (OFT). We observed a large range of stress-induced anxiety-like behaviors across the mouse lines (Fig. 1b; One-Way ANOVA, $F_{(15,198)}$ =2.262, p<0.001). We also observed an interaction between sex and genetic background (Two-Way ANOVA, $F_{(15,198)}$ =2.018, p<0.05). We then calculated the distance traveled in the open-field chamber and further observed a range of locomotor activity across mouse lines and sex (ANOVA, One-Way ANOVA, $F_{(15,224)}$ =19.38, p<0.001; Two-Way ANOVA, interaction $F_{(7,224)}$ =6.482, p<0.001).

Observing sex differences in response to CSDS amongst the different BXD lines tested, we examined the behavioral responses to CSDS of male mice independently from female mice (Fig. 1c,d). We observed that male DBA/2J and BXD22 were more susceptible to CSDS-induced social avoidance than the male founder C57BL/6J line (Kruskal–Wallis, $K_{(7,126)}$ =48.05; Dunns' corrected z=3.517, p=0.003; z=5.983, p<0.001). We also observed that male BXD5, 8, 22 and DBA/2J mice had lower distance traveled than the male founder C57BL/6J line (Kruskal–Wallis, $K_{(7,126)}$ =82.07; Dunns' corrected z=3.805, z=4.317, z=5.737, z=7.827, p<0.001).

Confirming the higher sensitivity of the DBA/2J mice to CSDS-induced social avoidance, female DBA/2J mice had a lower social interaction ratio than C57BL/6J female mice following CSDS (Fig. 1d, One-Way ANOVA, $F_{(7,105)=}2.213$; Bonferroni corrected t=3.156, p=0.01). While BXD14 male mice presented a similar degree of social interaction as male C57BL/6J line, female BXD14 had a lower social interaction ratio than female C57BL/6J mice (One-Way ANOVA, $F_{(7,105)=}2.213$; Bonferroni corrected *t*-tests, t=2.983, p=0.02). Additionally, we also observed that female DBA/2J and BXD22 (Kruskal–Wallis, K_(7,114)=33.78; Dunns' corrected z=4.510, p<0.001, z=2.771, p=0.04) lines had lower distances traveled than the female founder C57BL/6J line. Together, these results suggest that CSDS-induced behavioral outcomes are impacted by sex and genetic background.

C57BL/6J, DBA/2J, and BXD strains have distinct behavioral performance following nonsocial chronic stress

Adverse events are highly heterogeneous, and behavioral and physiological responses to stress vary with the nature of stressors. To capture the impact of genetics and sex on anxiety-like and exploratory behavior following chronic non-social stress exposure, we tested behavioral responses of BXD and founder C57BL/6J and DBA/2J mouse lines to the CVS paradigm. We and others have established that exposure to this non-social chronic stress disrupts exploratory behaviors, reward responses, and motivated behaviors^{34,36}.

We first exposed male and female mice from the BXD founder lines C57BL/6J and DBA/2J and the BXD8, BXD22, and BXD29 lines to the CVS paradigm (Fig. 2a). We then assessed anxiogenic behaviors in the novelty-suppressed feeding (NSF) task and observed a large range of behaviors across mouse lines and sexes (Fig. 2b; One-Way ANOVA, $F_{(9,141)}$ =9.474, p<0.001; see Supplementary Table S1 for details). Our results also showed distinct latency to feed between mouse lines and sex (Fig. 2b; Two-Way ANOVA, $F_{(4,141)}$ =17.18, p<0.001; Interaction $F_{(4,141)}$ =3.366, p=0.0072). We then tested the extent of CVS-induced anhedonia by measuring

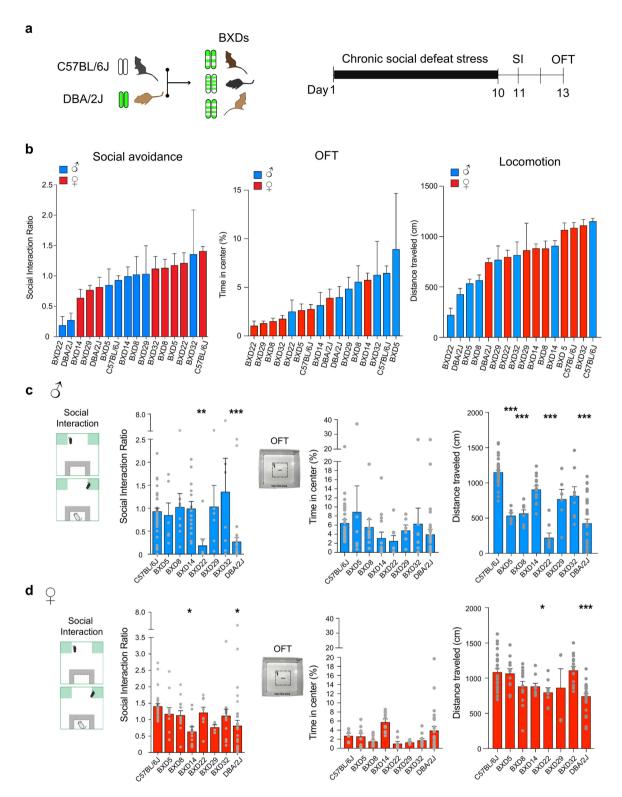


Fig. 1. C57BL/6J, DBA/2J, and BXD mice have distinct behaviors following chronic social defeat stress. (**a**) Experimental timeline illustrating the CSDS paradigm followed by the social interaction test (SI) and the open-field test (OFT). (**b**) Behavioral assessment showing the heterogeneous impact of CSDS on SI, OFT, and locomotion by genetic background in male (blue) and female (red) stressed mice. (**c**) CSDS-exposed male mice display distinct SI ratio, percentage of time spent in the center of the OFT, and distance traveled in the OFT in a "no-target" context amongst C57BL/6J, BXD: 5, 8, 14, 22, 29, and 32, and DBA/2J male mice following CSDS (n = 40, 6, 9, 14, 8, 6, 8, 35). (**d**) Same as c in CSDS-exposed female mice, C57BL/6J, BXD: 5, 8, 14, 22, 29, and 32, and DBA/2J (n = 28, 11, 12, 10, 8, 4, 14, 26). Bars represent mean ± SEM. ANOVA or Kruskal–Wallis tests followed by post-hoc comparison to C57BL/6J mice with Bonferroni correction; see Supplementary Table S1; *P < 0.05, **P < 0.01, and ***P < 0.01.

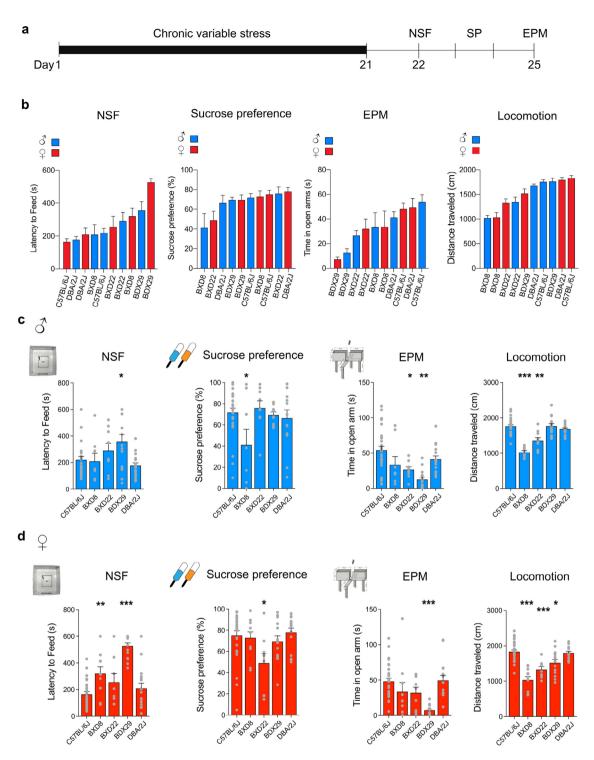


Fig. 2. C57BL/6J, DBA/2J, and BXD mice have distinct sensitivity to non-social chronic stress. (**a**) Experimental timeline illustrating the chronic variable stress (CVS) paradigm followed by the novelty suppress feeding test (NSF), the sucrose preference test (SP), the elevated plus-maze test (EPM), and locomotion in an open field. (**b**) Behavioral assessment showing the heterogeneous impact of CVS on NSF, SP, EPM, and locomotion by genetic background. Bars represent mean \pm SEM for n = 6–40 mice. (**c**) CVS-exposed male mice display distinct latency to feed during the NSF task, SP ratio (%), time spent as percentage in open arms of the EPM, and distance traveled in an open field in the "no-target" context amongst C57BL/6J, BXD: 8, 22, and 29 and DBA/2J male mice (n = 27, 8, 9, 15, 16). (**d**) CVS-exposed female mice display distinct latency to feed times during the NSF task, SP ratio (%), time spent and distance traveled in an open field in EPM open arms, and distance traveled in an open field in "no-target" context amongst C57BL/6J, BXD: 8, 22, and 29 and DBA/2J male mice (n = 27, 8, 9, 15, 16). (**d**) CVS-exposed female mice display distinct latency to feed times during the NSF task, SP ratio (%), time spent in EPM open arms, and distance traveled in an open field in "no-target" context amongst C57BL/6J, BXD: 8, 22, and 29 and DBA/2J (n = 27, 10, 9, 15, 15). Bars represent mean \pm SEM. ANOVA followed by post hoc comparison to C57BL/6 J mice with Bonferroni correction; see Supplementary Table S1; *P < 0.05, **P < 0.01.

the preference for sucrose solution over water (One-Way ANOVA, $F_{(9,133)} = 2.74$, p = 0.006). We observed an interacting effect of sex and genetics on sucrose preference (Two-Way ANOVA, $F_{(4,141)} = 3.998$, p = 0.004). Following these measurements, we tested exploratory behavior in an elevated plus maze (EPM, Fig. 2b; One-Way ANOVA, $F_{(9,140)} = 6.244$, p < 0.001). Further confirming the variable sensitivity to stressful stimuli, we observed a large range of locomotor behaviors across mouse lines and sex (Fig. 2b; One-Way ANOVA, $F_{(9,140)} = 6.244$, p < 0.001). Further confirming the variable sensitivity to stressful stimuli, we observed a large range of locomotor behaviors across mouse lines and sex (Fig. 2b; One-Way ANOVA, $F_{(9,142)} = 18.88$, p < 0.001; Two-Way ANOVA, gene $F_{(4,142)} = 39.61$, p < 0.0001, interaction $F_{(4,142)} = 2.368$, p = 0.0475). We observe that BXD29 had lower performances than the male founder C57BL/6J line on NSF (Fig. 2c; One-

We observe that BXD29 had lower performances than the male founder C57BL/6J line on NSF (Fig. 2c; One-Way ANOVA, $F_{(4,70)} = 3.304$, t = 2.783, p = 0.03) and CVS exposed BXD8 mice had a blunted sucrose preference (One-Way ANOVA, $F_{(4,62)} = 2.749$, t = 3.07, p = 0.01). Additionally, BXD22 and BXD29 males had higher anxiety levels when compared to the male founder C57BL/6J line (One-Way ANOVA, $F_{(4,70)} = 7.596$, *t*-tests, t = 2.915, p = 0.02; t = 5.272, p = 0.001; Fig. 2c). We also observed that the male BXD 8 and BXD22 (One-Way ANOVA, $F_{(4,70)} = 23.19$; *t*-tests, t = 8.39, t = 4.87, p < 0.001) lines had lower distances traveled than the male founder C57BL/6J line (Fig. 2c).

Confirming the heterogeneous stress-induced behavioral responses between sex and strain, we observed that female BXD8 and BXD29 mice displayed a higher latency to feed when compared to C57BL/6J female mice (Fig. 2d, One-Way ANOVA, $F_{(4,71)}$ =20.18, t=3.191, t=8.660, p=0.008, p<0.001). While sucrose preference in BXD22 males was similar to C57BL/6J males, BXD22 females had a lower sucrose preference ratio relative to C57BL/6J females (One-Way ANOVA, $F_{(4,71)}$ =3.155, *t*-tests, t=3.224, p=0.008). In addition to high NSF, BXD29 females displayed higher CVS-induced anxiety-like levels when compared to C57BL/6J females (One-Way ANOVA, $F_{(4,70)}$ =6.534, *t*-tests, t=4.677, p<0.001). We also observed that BXD8, BXD22, and BXD29 (One-Way ANOVA, $F_{(4,72)}$ =20.09, *t*-tests, t=7.987, t=4.673, t=3.359, p<0.001, p=0.005) females had lower distances traveled relative to C57BL/6J females. Together these results confirm that sex and strain have a relevant influence on chronic social and non-social stress behavioral responses.

Distinct morphine-induced motor effects between C57BL/6J and DBA/2J mouse lines following chronic stress exposure

Variability in response to psychoactive drugs in mice is known to depend in part on genetic differences^{37–40}. To test if genetics and sex contribute to the impact of chronic stress on morphine sensitivity, we measured morphine-induced motor activity in stressed male and female C57BL/6 J and DBA/2 J mice.

Following the CSDS or CVS paradigm, male and female C57BL/6J and DBA/2J mice were injected with 7.5 mg/kg of morphine (intraperitoneal; Fig. 3a). This dose was selected because it consistently and reproducibly produces a sub-maximal degree of behavioral responses in C57BL/6J mice (males and females), making it possible to detect manipulations that either increase or decrease sensitivity to the place conditioning effects of the drug. Fifteen minutes after the morphine injection, mice were placed in a large cage allowing for tracking of their locomotor activity. We observed that male and female C57BL/6J mice exposed to CSDS had higher locomotor activity when compared to DBA/2J mice (Fig. 3b; *t*-tests, t = 6.452, p < 0.001; t = 6.960, p < 0.001). Recapitulating these data, male and female C57BL/6 J mice exposed to CVS had higher locomotor activity when compared to DBA/2 mice (Fig. 3b; *t*-tests, t = 6.452, p < 0.001; t = 6.960, p < 0.001). Recapitulating

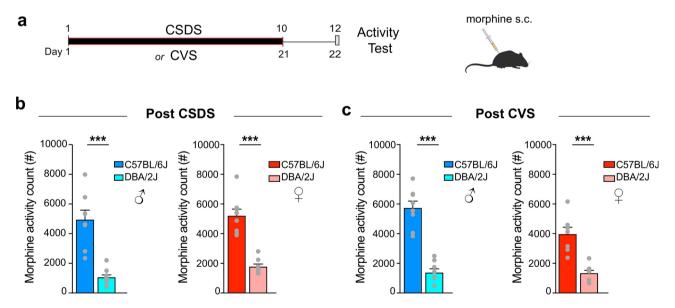


Fig. 3. Parental C57BL/6J and DBA/2J mice have distinct locomotor responses to acute morphine after chronic stress exposure. (**a**) Experimental timeline illustrating the two procedures performed, i.e. CSDS or CVS, followed by locomotor activity monitoring for 60 min in a locomotor apparatus 15 min after the animals were given an intraperitoneal injection of 7.5 mg/kg morphine. (**b**) Locomotor activity counts in C57BL/6J and DBA/2J male (left) and female (right) mice (n=8) exposed to CSDS. (**c**) Locomotor activity counts in C57BL/6J and DBA/2J male (left) and female (right) mice (n=8) exposed to CVS. Bars represent mean \pm SEM. Unpaired *t*-test; see Supplementary Table S1; *P < 0.05, **P < 0.01, and ***P < 0.01.

DBA/2J mice (Fig. 3c; *t*-tests, t=8.080, p<0.001; t=5.301, p<0.001). We observed that sex had an interacting effect with the mice's genotype on morphine-induced locomotor activity (Two-Way ANOVA, sex $F_{(1,27)}$ =6.025, p=0.02; interaction $F_{(1,27)}$ =5.517, p=0.02; gene effect $F_{(1,27)}$ =90.17, p<0.001; see Supplementary Table S1). However, we did not detect distinct behavioral responses of the C57BL/6j and DBA2J mice due to the different nature of the stress (Two-Way ANOVA, interaction $F_{(1,61)}$ =0.059, p=0.807; stress $F_{(1,61)}$ =0.174, p=0.736; gene effect $F_{(1,61)}$ =158.5, p<0.001; see Supplementary Table S1). Together, our results confirmed that the interaction between chronic stress and behavioral responses to morphine vary between genetic backgrounds.

Sex-specific impact on morphine's rewarding effects in stressed C57BL/6J and DBA/2J mice

To test the hypothesis that the rewarding properties of morphine differ between stressed C57BL/6J and DBA/2J male and female mice, we employed the CPP paradigm (Fig. 4a). Following CSDS or CVS, male and female C57BL/6J and DBA/2J mice were conditioned with once-daily alternating injections of saline and morphine (7.5 mg/kg, *s.c.*) over a period of 2 days. Side preference was assessed one day prior to morphine conditioning (pretest), and an unbiased CPP approach was utilized. One day after conditioning, the preference for the morphine-paired compartment was assessed.

We observed that CPP scores and preferences did not show significant differences between male CSDS-exposed C57BL/6J and DBA/2J mice (Fig. 4b; Two-Way ANOVA, $F_{(2,42)}=0.038$, p=0.84; *t*-test, t=0.315, p=0.75). Two-Way ANOVA comparison showed that, while CSDS-exposed female C57BL/6J mice established morphine CPP, CSDS-exposed female DBA/2J mice did not establish a preference for the morphine-paired compartment (Fig. 4b; Two-Way ANOVA, interaction $F_{(2,42)}=18.09$, p<0.0001; paired-side t=4.886 p<0.001, unpaired-side t=4.047 p=0.007). We further observed significant differences in preference for the morphine-paired compartment between CSDS female C57BL/6J and DBA/2J mice (Fig. 4c; *t*-test, t=3.677, p=0.002). We defined that sex had an effect on morphine CPP (Two-Way ANOVA, sex $F_{(1,28)}=3.995$, p=0.049; see Supplementary Table S1). Recapitulating these results, CVS-exposed female C57BL/6J and DBA/2J mice showed trending differing CPP profiles (Fig. 4d; Two-Way ANOVA, interaction $F_{(2,42)}=3.72$, p=0.059), whereas CVS-exposed male C57BL/6J and DBA/2J mice did not (Fig. 4e; Two-Way ANOVA, interaction $F_{(2,42)}=1.079$, p=0.35). Additionally, CVS-exposed male C57BL/6J and DBA/2J mice did not show a significant difference in preference for the morphine-paired side (Fig. 4d; *t*-test, t=0.791, p=0.44), and CVS-exposed female C57BL/6J mice established CPP, while DBA/2J females did not (Fig. 4e; *t*-test, t=2.677, p=0.047). We did not detect distinct behavioral responses of the C57BL/6j and DBA2J mice due to the different nature of the stress (Two-Way ANOVA, interaction $F_{(1,60)}=0.651$, p=0.42; stress $F_{(1,60)}=0.004$, p=0.94; gene effect $F_{(1,60)}=2.06$, p=0.15; see Supplementary Table S1). Our results establish a strain- and sex-specific impact on the rewarding properties of morphine in stressed C57BL/6J and DBA/2J mice.

Discussion

While stress is a well-known risk factor in the development of drug addiction, the genetic factors that make certain individuals particularly susceptible or resilient to stress and, thereby, more or less vulnerable to becoming addicted, remain elusive. In this study, we tested if genetic components map onto the behavioral and physiological mechanisms underlying (1) behavioral deficits following chronic stress exposure and (2) the ability of prior stress experience to influence drug responses. The BXD family of recombinant inbred strains derived from crossing two inbred parental mouse lines-C57BL/6J and DBA/2J-have been extensively used for almost 50 years in fields such as neuropharmacology²⁶, immunology⁴¹, and addiction^{18,25,38} to answer important biological questions. Combining the use of the BXD mouse lines and their founder lines with models of chronic stress exposure and morphine sensitivity, we first showed that CSDS-induced behavioral outcomes are impacted by sex and strain. Further, we established that vulnerability to chronic non-social stress (i.e. CVS) also varies depending on sex and genetics. We revealed that DBA/2J and BXD22 male and female mice are more susceptible to chronic social stress than C57BL/6J mice, evidenced by stronger social avoidance and anxiety-like behaviors. We observed sexual dimorphism in responses to CSDS amongst the BXD5, BXD8, BXD14, and BXD32 lines. To investigate the interaction between genetics and vulnerability to prolonged exposure to non-social stressors, we exposed C57BL/6J, DBA/2J, BXD8, BXD22, and BXD29 male and female mice to CVS and observed that DBA/2J female mice are more sensitive to CVS when compared to C57BL/6J female mice (i.e. a heightened decrease in sucrose preference). Confirming the stress vulnerability of BXD22 mice observed after CSDS, CVS-exposed male and female BXD22 mice displayed higher levels of anxiety-like measures than C57BL/6J mice. Interestingly, while BXD29 mice behaved like C57BL/6J after CSDS, both BXD29 female and male mice developed a higher anxiety profile following CVS when compared to C57BL/6J mice. Finally, we identified that DBA/2J and C57BL/6J mice pre-exposed to CSDS displayed differences in morphine sensitivity.

Here, we aimed to define the impact of genetics in the expression of the behavioral outcomes of chronic stress exposure, including social and exploratory behaviors, measures of anxiety-like endpoints, as well as morphine sensitivity. While our study highlights the difference in response to stress-eliciting stimuli, it is important to consider that pre-existing behavioral differences may also contribute to the observed behavioral divergence across mouse lines⁴¹. For example, in the absence of stress exposure, DBA mice show decreased reciprocal social behavior in the resident intruder task and the three-chamber sociability test⁴². Further, previous work has shown that social approach/sniff, ultrasonic vocalizations, partner sniffing, anxiety-like, and aggressive behavior all vary between various BXD strains⁴³⁻⁴⁵, which would also impact stress vulnerability. These strainspecific behaviors are reflected in differences in brain morphology as well. A stereological assessment of lateral septum (LS) volume revealed wide variability among the BXD strains and their parental lines, which, given the role of the LS in social and drug reward, may mediate GxE impacts on stress- and morphine-sensitivity^{46,47}. We observed that DBA/2J mice exhibit stronger social avoidance behaviors when compared to C57BL/6J mice after CSDS. Interestingly, following CVS exposure, DBA/2J and C57BL/6J mice exhibited similar behavioral

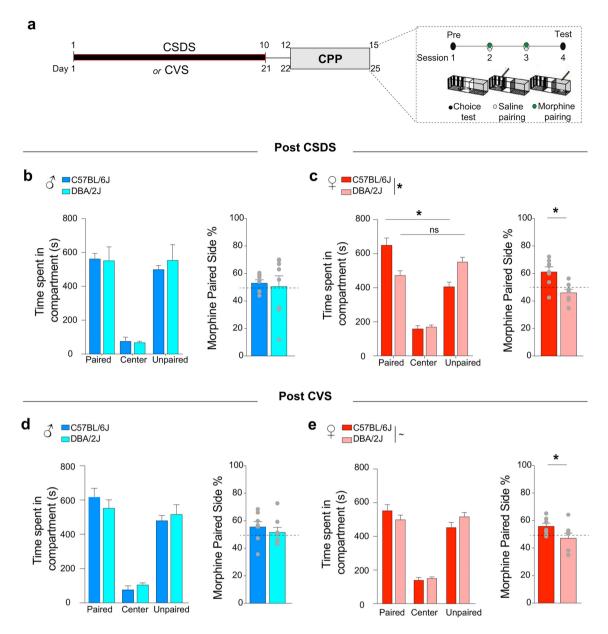


Fig. 4. Parental C57BL/6J and DBA/2J mice have distinct place conditioning responses to morphine after chronic stress exposure. (**a**) Experimental timeline illustrating the two procedures performed, i.e. CSDS or CVS, prior to performing morphine conditioned place preference (CPP). Morphine CPP was performed using an unbiased approach: one compartment was paired with saline (0.3 mL, *s.c.*) and the other paired with morphine (7.5 mg/kg, *s.c.*). Sessions were 20 min long. (**b**) Time spent in each compartment of the three-chamber apparatus after paired-conditioning with morphine (vs. saline) in CSDS-exposed C57BL/6J and DBA/2J male mice (n = 8) and respective percentage in the morphine-paired chamber. (**c**) Same as (**b**), in CSDS-exposed C57BL/6J and DBA/2J female mice (n = 8). (**d**) Time spent in each compartment of the three-chamber apparatus after paired-conditioning with morphine (vs. saline) in CVS-exposed C57BL/6J and DBA/2J female mice (n = 8). (**d**) Time spent in each compartment of the three-chamber apparatus after paired-conditioning with morphine (vs. saline) in CVS-exposed C57BL/6J and DBA/2J female mice (n = 8). (**d**) Time spent in each compartment of the three-chamber apparatus after paired-conditioning with morphine (vs. saline) in CVS-exposed C57BL/6J and DBA/2J female mice (n = 8). (**d**) Time spent in each compartment of the three-chamber apparatus after paired-conditioning with morphine (vs. saline) in CVS-exposed C57BL/6J and DBA/2J female mice (n = 8). Bars represent mean ± SEM. Two-way ANOVA followed by post-hoc comparison to C57BL/6J mice with Bonferroni correction and non-paired *t*-test; see Supplementary Table S1; *P < 0.05, **P < 0.01, and ***P < 0.01.

.....

responses. Conversely, while BXD29 mice behaved like C57BL/6J mice after CSDS, the BXD29 mice developed a heightened anxiety profile following CVS compared to C57BL/6J mice. These results suggest distinct genetic contributions in the mechanisms underlying these two stress-induced behavioral outcomes. In line with these observations, several studies have reported distinct and even opposite modifications of neural circuits involved in social and non-social stress exposure, including within the dopamine system^{32,48,49}. In particular, it has been shown that CSDS induces a hyper-dopaminergic state in mice expressing social avoidance and decreased sucrose preference. Oppositely, chronic non-social stress reduces the activity of midbrain dopaminergic neurons in mice

exhibiting depressive-like behaviors such as a lower sucrose preference ratio. Together, these results emphasize that opposite neuronal modifications, supported by distinct genetic mechanisms, may converge to similar pathological behavioral responses to distinct types of stressors.

In line with evidence in humans, we observed sexual dimorphism across the parental and BXD mouse lines in behavioral outcomes after chronic stress exposure. For example, we observed that, while male BXD22 mice developed stronger social avoidance behaviors compared to male C57BL/6J mice, female BXD22 mice did not and maintained social behaviors similar to those of female C57BL/6J mice exposed to CSDS. Notably, we observed that CSDS-exposed female mice had lower exploratory behaviors in the center of the open-field when compared to their male counterparts. Interestingly, as a result of prolonged exposure to non-social stressors, we observed similar behavioral alterations between male and female mice within the mouse lines. Both male and female BXD29 mice exhibited the strongest stress-induced behavioral deficits compared to C57BL/6J mice. These results indicate stress-specific sex differences in the mechanisms underlying behavioral responses to these two types of stress. In a broader assessment of fear-related behavior in the absence of stress, one study showed that anxiety-like behavior and fear-conditioning tendencies were independent of sensory and motor functions suggesting that these genetically divergent characteristics could also contribute to innate behavioral differences precipitated by chronic stress exposure⁵⁰.

Similar to the behavioral changes after chronic stress exposure⁵¹, sex is a key factor that may play a role in the development of addiction. Women tend to progress more rapidly than men through the stages of substance use disorder despite initiating drug use at a later age⁵². Women are more likely to report misusing prescription opioids to cope with negative affect, while also showing the characteristic rapid progression from first drug experience to developing substance use disorder^{53,54}. To allow for the detection of subtle but robust sex differences in behavioral responses to morphine, we utilized a threshold dose of morphine (7.5 mg/kg), below the dose known to induce CPP or somatic withdrawal in stress-naïve mice⁵⁵. We observed a robust strain effect on morphine-induced locomotor activity between the C57BL/6J and DBA/2J mice. Additionally, female C57BL/6J mice showed greater CPP when compared to female DBA/2J mice, while male DBA/2J and C57BL/6J mice did not establish morphine CPP at the dose used. We found that strain and sex contributed to stress-induced sensitivity to morphine reward, independently from the nature of the chronic stressor, i.e., social or non-social. Further genetic analyses seeking to identify the specific genetic loci involved in the impact of stress on morphine sensitivity will help identify novel mechanisms in gene x environment interactions in neuropsychiatric disorders.

To conclude, our results support the hypothesis that genetic variations contribute to stress responses, such as social avoidance and anxiety-like behaviors, and influence sensitivity to morphine and, as a result, presumably modulate the risk of psychiatric disorders and addiction. Our previous studies performed in isogenic mouse lines have allowed the identification of physiological processes and epigenetic mechanisms contributing to stress-susceptibility and drug responses on a single genetic background^{56–58}. In parallel, important genetic mapping research efforts using recombinant inbred mouse models have identified the high degree of genetic homology between humans and mice⁵⁹. This homology has allowed for impactful cross-species studies that will define the genes interacting in the regulation of behavioral stress responses and addictive behaviors. Characterization of the genetic, neurobiological, social, and environmental factors that mediate addiction risk will fundamentally improve our understanding of individual variations in psychiatric disorders and responses to drugs of abuse and provide highly useful information for the development of new treatment strategies and, eventually, prevention measures.

Material and methods

Mice

The study is reported in accordance with the Animal Research: Reporting of In Vivo Experiments (ARRIVE) guidelines (https://arriveguidelines.org). C57BL/6J, DBA2J, and BXD female and male mice (7–12 weeks old; Jackson Laboratory, Bar Harbor) were used for all experiments and habituated to the animal facility for one week before experimental manipulations. Mice were housed in groups of five at a constant ambient temperature $(24 \pm 1 \text{ °C})$ under a 12-h light/dark cycle (lights on from 7:00 A.M.) with ad libitum access to water and food, except when otherwise specified for behavioral testing. Following experiments, mice were euthanized with lethal doses of anesthesia (ketamine 300 mg/kg/xylazine 150 mg/kg) followed by cervical dislocation. All efforts were made to minimize the number of animals used in the experiments and to limit their distress and suffering. Experiments were conducted in accordance with the guidelines of the Institutional Animal Care and Use Committee (IACUC) at Mount Sinai. Experimental protocols were approved by the Institutional Animal Care and Use Committee (IACUC) at Mount Sinai.

Chronic social defeat stress (CSDS) paradigm

Pre-screened CD-1 retired male breeder mice were singled-housed in one compartment of a large mouse cage partitioned by a perforated Plexiglass divider vertically placed through the cage, which has a metal top to hold food and water on both sides. This is used as the resident cage and is the permanent home for the CD-1 mouse. The CSDS paradigm begins when an experimental mouse is introduced into the CD-1 home cage and the ensuing fighting is allowed to occur for 10 min. Following the interaction, mice are individually confined to one compartment for the rest of the day, allowing for continued sensory exposure. This is repeated for 10 days, and the effect of the 10-day defeat is measured on day 11 through a social interaction test³². Similar to the male CSDS paradigm, female mice are placed, as an intruder, in the home cage of an Esr1-Cre mouse. The Esr1-Cre mouse is injected with CNO, an exogenous drug targeting the DREADD expressed in the ventrolateral subdivision of the ventromedial hypothalamus, to induce aggressive behavior toward the female experimental mice. Following 5 min of physical contact, the female mice are placed back into their home-cage, where they are grouped housed. Similar to the male CSDS paradigm, this schedule is repeated for 10 days³³.

Social interaction test (SI test)

Mice were placed in an OFT box containing a wire mesh cage on one side under red-light conditions (<15 lux)³². The open field arena and wire-mesh enclosures were thoroughly cleaned between mice with an odorless 5% ethanol solution. The CSDS-exposed mice were individually placed in the open field box and allowed to explore for 2.5 min (without a social target present, known as the "no target phase"). The experimental mouse was then removed, the open field was cleaned, and an unfamiliar aggressor (CD-1/Esr1-Cre) mouse was placed into the mesh cage (known as the "target phase"). The experimental mouse was then allowed to explore for another 2.5 min. Time spent interacting with the social target and locomotion were measured using an automated video tracking system (Ethovision). Experimental mice were then placed back into their home cage (single-housed). The social interaction ratio was calculated as [time in social interaction zone with "no target"], as described previously^{32,60}. The distance traveled was analyzed during the "no target" phase to avoid potential biases due to the novel social target being present.

Chronic variable stress (CVS)

CVS consists of three different stressors over 30 days³⁴. Environmental stressors are presented on an alternating schedule to prevent habituation. Stressors are administered in the following order: 100 random mild foot shocks at 0.45 mA for one hour (10 mice to a chamber), a tail suspension stress for one hour, and restraint stress (placed inside a 50 mL falcon tube) for one hour. The three stressors are then repeated in the same cycle for the remainder of the stress protocol.

Open field test (OFT)

Mice were placed in the open field arena $(44 \times 44 \text{ cm})$ for 5 min to compare the distance traveled and time spent in the peripheral zone compared to the center zone $(10 \times 10 \text{ cm})$. Testing conditions occurred under red-light conditions (<10 lux) in a room isolated from external sound sources. The apparatus was thoroughly cleaned between mice with an odorless 5% ethanol solution. The mouse's time spent in specific open field areas—was video-tracked and scored with Ethovision software³².

Elevated plus maze test (EPM)

The EPM was designed in black Plexiglass (L/W/D: 70/5/20 cm) and fitted with white surfaces to provide contrast. Testing conditions occurred under red-light conditions (<10 lux) in a room isolated from external sound sources. The EPM apparatus was thoroughly cleaned between mice with an odorless 5% ethanol solution. Mice were positioned in the center of the maze, and behavior was video-tracked for 5 min³². Time in EPM compartments was measured using a video-tracking system (Ethovision) set to focus on the mouse center-point at the commencement of each trial.

Sucrose preference (SP)

Mice were habituated to having access to two water bottles (50-ml tubes with fitted ball-point sipper tubes) for one day, and were then given free access to both water and a 1% sucrose solution, for two consecutive days³². Bottles were weighed daily and interchanged (left to right, right to left) to avoid biases from a potential side preference. Sucrose preference scores were calculated as ([sucrose solution consumed]/[sucrose+water solutions consumed]) \times 100.

Novelty suppressed feeding (NSF)

The NSF test elicits conflicting motivations in mice: the drive to eat vs. the fear of moving to the center of a novel, open arena⁴⁹. Mice were food-restricted for 24 h prior to testing³⁴. Under red-light conditions, mice were placed in the corner of a clear plastic testing chamber $(50 \times 50 \times 20 \text{ cm})$ lined with corncob bedding. A single food pellet was present in the center of the box. The latency to begin consumption in a 6-min test was measured.

Morphine conditioned place preference (CPP)

Morphine CPP was conducted as previously described⁵⁸. We used an unbiased, three-compartment apparatus. First day 0, mice were allowed to freely explore the entire apparatus for 30 min to obtain baseline preference to any of the three compartments; we know that mice show no inherent group biases for a given chamber across multiple strains⁵². Mice were then given conditioning trials (two per day) on two consecutive days. For both trials, mice received saline (0.3 mL, subcutaneous; *s.c.*) and were confined to one of the compartments of the apparatus. After 4 h, mice received morphine (7.5 mg/kg in 0.3 mL, *s.c.*) and were confined to the opposite compartment. The drug-paired chamber is randomized across mice. On test day (day 4), mice are again allowed to freely explore the entire apparatus for 30 min. CPP scores are calculated as the time spent in the morphine-paired chamber minus the time spent in the saline-paired chamber. The time spent in each compartment was determined using an automated system and CPP scores were calculated as the percent of time spent in the morphine-paired compartment.

Locomotor sensitization

Locomotor activity was individually monitored in specialized locomotor chambers with no bedding, 15 min after the animals were given an intraperitoneal injection of 7.5 mg/kg morphine. Locomotor activity was monitored for 60 min.

Data analyses and statistics

All behaviors were monitored and scored using automated and unbiased Ethovision software^{32,51}. Experimenters analyzing the dataset were blinded to the experimental conditions. The statistical analyses were performed using

Graphpad Prism (version 8, La Jolla, CA, USA) and R (version 3.3.3) software and collected in Supplementary Table S1. The normality of the distributions was assessed using Kolmogorov–Smirnov tests. The statistical analyses were performed considering the sample size, normality, and homoscedasticity of the distributions. The data fitting assumptions of the general linear model were subjected to two-sided Student's *t*-tests, or multiple comparisons using a one-way, two-way, or repeated measures (RM) ANOVA followed by post hoc two-sided *t*-tests with Bonferroni correction for multiple comparisons (*t*-test, p-values). Non-parametric Kruskal–Wallis followed by post hoc Dunn's corrected comparisons and Mann–Whitney two-sided statistical analyses were performed for datasets that did not follow a normal distribution. The statistical significance threshold was set at 0.05.

Data availability

All data generated and analyzed during this study are included in this article and available in Supplementary Table S2.

Received: 23 February 2024; Accepted: 21 November 2024 Published online: 28 December 2024

References

- 1. Shurtleff, D., Sasek, C. & Kautz, M. Sponsor's foreword: NIDA at forty. Neuropharmacology 76(Pt B), 195-197 (2014).
- Kendler, K. S., Karkowski, L. M., Neale, M. C. & Prescott, C. A. Illicit psychoactive substance use, heavy use, abuse, and dependence in a US population-based sample of male twins. Arch. Gen. Psychiatry 57, 261–269 (2000).
- 3. Tsuang, M. T., Bar, J. L., Harley, R. M. & Lyons, M. J. The harvard twin study of substance abuse: what we have learned. *Harv. Rev. Psychiatry* 9, 267–279 (2001).
- 4. Carlezon, W. A. J., Duman, R. S. & Nestler, E. J. The many faces of CREB. Trends Neurosci. 28, 436-445 (2005).
- Sinha, R. Chronic stress, drug use, and vulnerability to addiction. Ann. N. Y. Acad. Sci. 1141, 105–130 (2008).
 Khibnik, L. A. et al. Stress and cocaine trigger divergent and cell type-specific regulation of synaptic transmission at single spines
- in nucleus accumbens. *Biol. Psychiatry* 79, 898–905 (2016).
 7. Morel, C. et al. Nicotinic receptors mediate stress-nicotine detrimental interplay via dopamine cells' activity. *Mol. Psychiatry* 23, 1597–1605 (2018).
- 8. Andersen, S. L. Stress, sensitive periods, and substance abuse. Neurobiol. Stress 10, 100140 (2019).
- Crabb, D. W., Edenberg, H. J., Bosron, W. F. & Li, T. K. Genotypes for aldehyde dehydrogenase deficiency and alcohol sensitivity. The inactive ALDH2(2) allele is dominant. J. Clin. Investig. 83, 314–316 (1989).
- 10. Burton, P. R., Tobin, M. D. & Hopper, J. L. Key concepts in genetic epidemiology. Lancet (London, England) 366, 941-951 (2005).
- Kreek, M. J., Nielsen, D. A. & LaForge, K. S. Genes associated with addiction: alcoholism, opiate, and cocaine addiction. *Neuromol. Med.* 5, 85–108 (2004).
- Deak, J. D. et al. Genome-wide association study in individuals of European and African ancestry and multi-trait analysis of opioid use disorder identifies 19 independent genome-wide significant risk loci. *Mol. Psychiatry* 27, 3970–3979 (2022).
- Hatoum, A. S. et al. Multivariate genome-wide association meta-analysis of over 1 million subjects identifies loci underlying multiple substance use disorders. Nat. Ment. Health 1, 210–223 (2023).
- 14. Strain, E. C. Assessment and treatment of comorbid psychiatric disorders in opioid-dependent patients. Clin. J. Pain 18, S14-27 (2002).
- Conway, K. P., Compton, W., Stinson, F. S. & Grant, B. F. Lifetime comorbidity of DSM-IV mood and anxiety disorders and specific drug use disorders: results from the National Epidemiologic Survey on Alcohol and Related Conditions. J. Clin. Psychiatry 67, 247–257 (2006).
- Hser, Y.-I. et al. Long-term follow-up assessment of opioid use outcomes among individuals with comorbid mental disorders and opioid use disorder treated with buprenorphine or methadone in a randomized clinical trial. Addiction 117, 151–161 (2022).
- 17. Schenk, S., Eisenbarth, H. & Dixon, L. Treating opioid use disorders in the criminal justice system with pharmacotherapy. *Forensic Sci. Int. Mind Law* 1, 100009 (2020).
- Ashbrook, D. G. et al. A platform for experimental precision medicine: The extended BXD mouse family. *Cell Syst.* 12, 235-247.e9 (2021).
- Williams, R. W., Strom, R. C., Rice, D. S. & Goldowitz, D. Genetic and environmental control of variation in retinal ganglion cell number in mice. J. Neurosci. Off. J. Soc. Neurosci. 16, 7193–7205 (1996).
- Williams, R. W., Gu, J., Qi, S. & Lu, L. The genetic structure of recombinant inbred mice: high-resolution consensus maps for complex trait analysis. *Genome Biol.* 2, RESEARCH0046 (2001).
- Andreux, P. A. et al. Systems genetics of metabolism: the use of the BXD murine reference panel for multiscalar integration of traits. Cell 150, 1287-1299 (2012).
- 22. McKnite, A. M. et al. Murine gut microbiota is defined by host genetics and modulates variation of metabolic traits. *PLoS One* 7, e39191 (2012).
- 23. Bachmanov, A. A. et al. Genetics of sweet taste preferences. Flavour Fragr. J. 26, 286-294 (2011).
- 24. Ashbrook, D. G., Williams, R. W., Lu, L. & Hager, R. A cross-species genetic analysis identifies candidate genes for mouse anxiety and human bipolar disorder. *Front. Behav. Neurosci.* 9, 171 (2015).
- Chunduri, A., Watson, P. M. & Ashbrook, D. G. New insights on gene by environmental effects of drugs of abuse in animal models using GeneNetwork. *Genes (Basel)* 13, 614 (2022).
- Grisel, J. E. et al. Quantitative trait loci affecting methamphetamine responses in BXD recombinant inbred mouse strains. J. Neurosci. Off. J. Soc. Neurosci. 17, 745–754 (1997).
- Crabbe, J. C., Kosobud, A., Young, E. R. & Janowsky, J. S. Polygenic and single-gene determination of responses to ethanol in BXD/ Ty recombinant inbred mouse strains. *Neurobehav. Toxicol. Teratol.* 5, 181–187 (1983).
- Tolliver, B. K., Belknap, J. K., Woods, W. E. & Carney, J. M. Genetic analysis of sensitization and tolerance to cocaine. J. Pharmacol. Exp. Ther. 270, 1230–1238 (1994).
- 29. Wahlström, A., Hammar, L., Lundin, L. G. & Rane, A. Morphine metabolism in mouse brain. *NIDA Res. Monogr.* **75**, 603–606 (1986).
- Phillips, T. J., Belknap, J. K. & Crabbe, J. C. Use of recombinant inbred strains to assess vulnerability to drug abuse at the genetic level. J. Addict. Dis. 10, 73–87 (1991).
- Badea, A., Johnson, G. A. & Williams, R. W. Genetic dissection of the mouse CNS using magnetic resonance microscopy. *Curr. Opin. Neurol.* 22, 379–386 (2009).
- 32. Morel, C. et al. Midbrain projection to the basolateral amygdala encodes anxiety-like but not depression-like behaviors. *Nat. Commun.* **13**, 1532 (2022).

- Golden, S. A., Covington, H. E., Berton, O. & Russo, S. J. A standardized protocol for repeated social defeat stress in mice. *Nat. Protoc.* 6, 1183–1191 (2011).
- 34. Hodes, G. E. et al. Sex differences in nucleus accumbens transcriptome profiles associated with susceptibility versus resilience to subchronic variable stress. J. Neurosci. 35, 16362–16376 (2015).
- 35. Barik, J. et al. Chronic stress triggers social aversion via glucococorticoid receptor in dopaminoceptive neurons. *Science* 339, 332-335 (2013).
- 36. Sinha, R. How does stress increase risk of drug abuse and relapse?. Psychopharmacology (Berl.) 158, 343-359 (2001).
- 37. Crabbe, J. C. & Belknap, J. K. Behavior genetic analyses of drug withdrawal. Alcohol Alcohol Suppl. 2, 477-482 (1993).
- 38. Baker, J. A., Brettin, J. T., Mulligan, M. K. & Hamre, K. M. Effects of genetics and sex on acute gene expression changes in the hippocampus following neonatal ethanol exposure in BXD recombinant inbred mouse strains. *Brain Sci.* **12**, 1634 (2022).
- Kitahama, K. & Valatx, J. L. Strain differences in amphetamine sensitivity in mice. II. Overcompensation of paradoxical sleep after deprivation in two C57 strains. *Psychopharmacology (Berl.)* 66, 291–295 (1979).
- Belknap, J. K. et al. Localization to chromosome 10 of a locus influencing morphine analgesia in crosses derived from C57BL/6 and DBA/2 strains. *Life Sci.* 57, PL117-24 (1995).
- Grizzle, W. E. et al. BXD recombinant inbred mice represent a novel T cell-mediated immune response tumor model. Int. J. Cancer 101, 270–279 (2002).
- Mulligan, M. K., Mozhui, K., Prins, P. & Williams, R. W. GeneNetwork: A toolbox for systems genetics. *Methods Mol. Biol.* 1488, 75–120. https://doi.org/10.1007/978-1-4939-6427-7_4 (2017).
- Ma, L., Piirainen, S., Kulesskaya, N., Rauvala, H. & Tian, L. Association of brain immune genes with social behavior of inbred mouse strains. J. Neuroinflamm. 18(12), 75. https://doi.org/10.1186/s12974-015-0297-5 (2015).
- Knoll, A. T., Jiang, K. & Levitt, P. Quantitative trait locus mapping and analysis of heritable variation in affiliative social behavior and co-occurring traits. *Genes Brain Behav.* 17(5), e12431. https://doi.org/10.1111/gbb.12431 (2018).
- 45. Delprato, A. et al. A quantitative trait locus on chromosome 1 modulates intermale aggression in mice. *Genes Brain Behav.* **17**(7), e12469. https://doi.org/10.1111/gbb.12469 (2018).
- 46. Talishinsky, A. & Rosen, G. D. Systems genetics of the lateral septal nucleus in mouse: heritability, genetic control, and covariation with behavioral and morphological traits. *PLoS One* 7(8), e44236. https://doi.org/10.1371/journal.pone.0044236 (2012).
- ZióŁkowska, B., Gieryk, A., Solecki, W. & PrzewŁocki, R. Temporal and anatomic patterns of immediate-early gene expression in the forebrain of C57BL/6 and DBA/2 mice after morphine administration. *Neuroscience* 22(284), 107–124. https://doi.org/10.101 6/j.neuroscience.2014.09.069 (2015).
- Friedman, A. K. et al. Enhancing depression mechanisms in midbrain dopamine neurons achieves homeostatic resilience. Science https://doi.org/10.1126/science.1249240 (2014).
- Zhang, S. et al. Sex Differences in the neuroadaptations of reward-related circuits in response to subchronic variable stress. Neuroscience https://doi.org/10.1016/j.neuroscience.2018.02.021 (2018).
- Brigman, J. L., Mathur, P., Lu, L., Williams, R. W. & Holmes, A. Genetic relationship between anxiety-related and fear-related behaviors in BXD recombinant inbred mice. *Behav. Pharmacol.* 20(2), 204–209. https://doi.org/10.1097/FBP.0b013e32830c368c (2009).
- 51. Li, L. et al. Social trauma engages lateral septum circuitry to occlude social reward. Nature 613, 696-703 (2023).
- 52. McHugh, R. K., Votaw, V. R., Sugarman, D. E. & Greenfield, S. F. Sex and gender differences in substance use disorders. *Clin. Psychol. Rev.* 66, 12–23 (2018).
- Jamison, R. N., Butler, S. F., Budman, S. H., Edwards, R. R. & Wasan, A. D. Gender differences in risk factors for aberrant prescription opioid use. J. Pain 11, 312–320 (2010).
- 54. Lewis, B., Hoffman, L. A. & Nixon, S. J. Sex differences in drug use among polysubstance users. *Drug Alcohol Depend.* 145, 127–133 (2014).
- 55. Papaleo, F. & Contarino, A. Gender- and morphine dose-linked expression of spontaneous somatic opiate withdrawal in mice. *Behav. Brain Res.* **170**, 110–118 (2006).
- 56. Kiraly, D. D. et al. Alterations of the host microbiome affect behavioral responses to cocaine. Sci. Rep. 6, 35455 (2016).
- 57. Issler, O. et al. The long noncoding RNA FEDORA is a cell type- and sex-specific regulator of depression. *Sci. Adv.* **8**, eabn9494 (2022).
- 58. Calipari, E. S. et al. Dopaminergic dynamics underlying sex-specific cocaine reward. Nat. Commun. 8, 13877 (2017).
- Emes, R. D., Goodstadt, L., Winter, E. E. & Ponting, C. P. Comparison of the genomes of human and mouse lays the foundation of genome zoology. *Hum. Mol. Genet.* 12, 701–709 (2003).
- 60. Takahashi, A. et al. Establishment of a repeated social defeat stress model in female mice. Sci. Rep. 7, 12838 (2017).

Acknowledgements

This work was supported by National Institute of Mental Health grants R01MH051399 (EJN) and R01MH120514 (CM, SJR) and by National Institute on Drug Abuse grant P01DA047233 (EJN), National Key R&D Program of China 2021ZD0202900 & 2021ZD0202902 (MHH), Research Fund for International Senior Scientists T2250710685 (MHH), Shenzhen Natural Science Foundation J20220127 (MHH) Shenzhen Medical Research Fund SMRF B2303012 (MHH), Shenzhen Key Laboratory of Precision Diagnosis and Treatment of Depression ZDSYS20220606100606014 (MHH), Science and Technology Research and Development Foundation of Shenzhen (High-level Talent Innovation and Entrepreneurship Plan of Shenzhen Team Funding) KQTD20221101093608028 (MHH). This study was also supported by a NARSAD Young Investigator Grant from the Brain & Behavior Research Foundation 29699 (CM) and 31194 (LFP), Leon Levy Foundation (LFP), the Alzheimer's Association (Grant No. 24AARG-NTF-1198746 to CM), and the Hope for Depression Research Foundation (EJN).

Author contributions

C.M., L.P., Y.V.Z., O.I., M.C., K.C., A.V.A., A.B., performed the behavioral assessments and with the assistance of C.B., C.M., L.P. Y.V.Z., M.C., O.I., analyzed the results. C.M., L.P., S.J.R., E.J.N., and M.H.H. designed the experiments with the assistance of R.W.W. M.K.M. C.M., L.P., O.I., S.J.R., E.J.N., and M.H.H. interpreted the results and wrote the paper, which was edited by all authors.

Declarations

Competing interests

The authors declare no competing interests.

Additional information

Supplementary Information The online version contains supplementary material available at https://doi.org/1 0.1038/s41598-024-80767-7.

Correspondence and requests for materials should be addressed to S.J.R., E.J.N. or M.-H.H.

Reprints and permissions information is available at www.nature.com/reprints.

Publisher's note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Open Access This article is licensed under a Creative Commons Attribution-NonCommercial-NoDerivatives 4.0 International License, which permits any non-commercial use, sharing, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if you modified the licensed material. You do not have permission under this licence to share adapted material derived from this article or parts of it. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by-nc-nd/4.0/.

© The Author(s) 2024