

MicroRNA-19b Associates with Ago2 in the Amygdala Following Chronic Stress and Regulates the Adrenergic Receptor Beta 1

Naama Volk,^{1,2} Evan D. Paul,^{2,3} Sharon Haramati,¹ Chen Eitan,¹ Brandon K.K. Fields,³ Raaya Zwang,¹ Shosh Gil,¹ Christopher A. Lowry,³ and Alon Chen^{1,2}

¹Department of Neurobiology, Weizmann Institute of Science, Rehovot 76100, Israel, ²Department of Stress Neurobiology and Neurogenetics, Max Planck Institute of Psychiatry, Munich 80804, Germany, and ³Department of Integrative Physiology, University of Colorado Boulder, Boulder, 80309 Colorado

Activation of the stress response in the presence of diverse challenges requires numerous adaptive molecular and cellular changes. To identify specific microRNA molecules that are altered following chronic stress, mice were subjected to the chronic social defeat procedure. The amygdala from these mice was collected and a screen for microRNAs that were recruited to the RNA-induced silencing complex and differentially expressed between the stressed and unstressed mice was conducted. One of the microRNAs that were significantly altered was microRNA-19b (miR-19b). Bioinformatics analysis revealed the adrenergic receptor β -1 (Adrb1) as a potential target for this microRNA with multiple conserved seed sites. Consistent with its putative regulation by miR-19b, Adrb1 levels were reduced in the basolateral amygdala (BLA) following chronic stress. *In vitro* studies using luciferase assays showed a direct effect of miR-19b on Adrb1 levels, which were not evident when miR-19b seed sequences at the Adrb1 transcript were mutated. To assess the role of miR-19b in memory stabilization, previously attributed to BLA-Adrb1, we constructed lentiviruses designed to overexpress or knockdown miR-19b. Interestingly, adult mice injected bilaterally with miR-19b into the BLA showed lower freezing time relative to control in the cue fear conditioning test, and deregulation of noradrenergic circuits, consistent with downregulation of Adrb1 levels. Knockdown of endogenous BLA-miR-19b levels resulted in opposite behavioral and noradrenergic profile with higher freezing time and increase 3-methoxy-4-hydroxyphenylglycol/noradrenaline ratio. These findings suggest a key role for miR-19b in modulating behavioral responses to chronic stress and Adrb1 as an important target of miR-19b in stress-linked brain regions.

Key words: adrenergic receptor beta 1; Ago2; amygdala; microRNA; stress

Introduction

The amygdala has a central role in regulating the behavioral stress response following stressful and emotional stimulation (Galvez et al., 1996). Amygdalar adrenergic receptor β 1 (Adrb1) was previously shown to affect anxiety-like behavior (Rudoy and Van Bockstaele, 2007; Fu et al., 2008) and fear memory (Roosendaal et al., 2004, 2006a). The Adrb1 is expressed in various stress-related

brain nuclei including the amygdala, hippocampus, and paraventricular nucleus of the hypothalamus (PVN). Infusion of β -adrenergic receptor antagonists into the amygdala blocks the modulation of memory storage by the stress-related hormones epinephrine and glucocorticoids (McGaugh et al., 1996), whereas drugs that enhance memory increase levels of noradrenaline in the amygdala (Hatfield and McGaugh, 1999). Interestingly, studies have convincingly demonstrated that the consolidation of aversive memories is facilitated by crosstalk between glucocorticoids, noradrenergic, and cannabinoid signaling in the basolateral nucleus of the amygdala (BLA; Roosendaal et al., 2006b). The regulation of the adrenergic receptor protein levels in the BLA is important for consolidation of emotionally aversive memories and coping with stressful challenges (Hill and McEwen, 2009). However, data regarding stress-dependent mechanisms mediating the protein levels of adrenergic receptors in the BLA is limited. Regulation at the transcript level of these receptors might enable a fast response during changing conditions and may be mediated by specific microRNA (miRNA) molecules.

miRNAs are short (~22 nucleotide), endogenous, single-stranded RNA molecules that regulate gene expression by promoting RNA transcript degradation or translation inhibition of target mRNAs. Several studies have reported that a range of be-

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Correspondence should be addressed to Dr Alon Chen, Department of Neurobiology, Weizmann Institute of Science, Rehovot, 76100, Israel. E-mail: alon.chen@weizmann.ac.il.

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havioral and pharmacological manipulations alters miRNA levels in adult mouse or rat brain structures (Uchida et al., 2008; Meerson et al., 2010; Lin et al., 2011; Issler et al., 2014). We previously reported an increase in several miRNAs including miR-34c in the amygdala 90 min following acute restraint stress (Haramati et al., 2011). miR-34c was found to regulate the stress-related corticotropin-releasing factor receptor type 1, and had a protective effect against stress-induced anxiety when lentivirally overexpressed in the mouse central amygdala. In addition, the brain enriched miRNA, miR-128b, was found to be upregulated following fear-extinction (Lin et al., 2011). Notably, changes in miRNA levels are not necessarily indicative for their immediate activity; it is only when a specific miRNA has matured, and has been incorporated into the RNA-induced silencing complex (RISC) in the presence of argonaute RISC catalytic component 2 (Ago2), where it associates with its target mRNA, that it is truly active (Meister et al., 2004).

To identify specific miRNAs that are activated (recruited to the RISC complex) following chronic stress, we have isolated and analyzed the Ago2-associated miRNAs in the amygdala of mice subjected to a social defeat protocol. One such prominent miRNA is miR-19b, which we further demonstrated both *in vitro* and *in vivo* to be a major regulator of *Adrb1* levels and related behavioral functions.

Materials and Methods

Animals

Male C57BL/6J and ICR (outbred mice strain, also known as CD1) mice (Harlan) were maintained in a pathogen-free temperature-controlled ($22 \pm 1^\circ\text{C}$) mouse facility on a reverse 12 h light/dark cycle at the Weizmann Institute of Science, according to institutional guidelines. Food (Harlan) and water were given *ad libitum*. The total number of animals used for the Ago2 IP was 30 (18 social defeat and 12 control). The total number of animal used for the lentiviruses injected mice was 40 in total (10/group).

Chronic social defeat

Ten-week-old C57BL/6J mice were subjected to a social defeat protocol as previously described (Krishnan et al., 2007). Briefly, the mice were placed in a homecage of an aggressive ICR mouse and allowed to physically interact for 5 min. During this time, the ICR mouse attacked the intruder mouse and the intruder displayed subordinate posturing. A perforated clear Plexiglas divider was then placed between the animals and the mice remained in the same cage for 24 h to allow sensory contact. The procedure was then repeated with an unfamiliar ICR mouse for each of the 10 consecutive days. Control mice were housed in the same room as the social defeat mice but were escorted out of the room during the 5 min interaction with the ICR. Control mice were handled daily and housed two in a cage with a perforated clear Plexiglas divider placed between the two mice.

Microdissection of brain sites for Ago2 IP

Amygdala samples were collected from social defeat and control mice 8 d after the end of the social defeat protocol. Tissue collection and processing was as previously described (Sztainberg et al., 2010; Lebow et al., 2012). Briefly, after removing the brain and placing it on an acryl 1 mm brain matrix (Stoelting, Catalog #51380), slices of 2 mm were taken using standard razor blades (GEM, 62-0165) based on designated anatomical markers. Blunted syringes at different diameters were used to extract the amygdala from slices removed from the matrix.

Immunoprecipitation of Ago2 protein

Pools of three amygdala from three mice from the same treatment group (either social defeat $n = 18$ or control $n = 12$) were homogenized in NP40 buffer, which was supplemented with RNase inhibitor, protease inhibitor and phosphatase inhibitor (Roche). The samples were constantly agitated for 2 h at 4°C . Samples were then centrifuged for 20 min

at 12,000 rpm at 4°C in a microcentrifuge; the supernatant was placed in a fresh tube kept on ice and the pellet was discarded. Magnetic protein G beads (Dynabeads, Invitrogen) were incubated with the Ago2 monoclonal antibody (WAKO) with rotation at room temperature for 10 min. After several washes, the samples were added to the Ago2-coated protein G beads and incubated overnight at 4°C under agitation. The following day the beads were washed three times with PBS. For RNA purification, the beads were homogenized in RLT buffer (RNeasy kit, miRNA supplementary protocol). For Western blot analysis, the beads were boiled in sample buffer to release the protein from the beads.

RNA purification and microarray

RNA from the Ago2 immunoprecipitation samples was isolated using the RNeasy plus kit (Qiagen) following Qiagen supplementary Protocol 1: purification of total RNA containing miRNA. RNA for all other purposes was isolated from frozen brain punches using miRNeasy mini kit (Qiagen) according to the manufacturer recommendation. RNA derived from tissues of stressed mice following Ago2 immunoprecipitation was further analyzed on Affymetrix miRNA 2.0 array (enriched RNA protocol).

Bioinformatic analysis of microRNA microarray results

miRNAs and genes were tested in 3 different web based programs in search for a seed match between a miRNA and a 3' UTR: Target Scan (<http://www.targetscan.org>), Miranda (<http://www.microrna.org>; Betel et al., 2008), and Pictar (<http://pictar.mdc-berlin.de>).

Cloning of 3' UTRs into *Psicheck2* luciferase expression plasmid

3' UTR sequence of *Adrb1* was PCR amplified from mouse genomic DNA using a forward primer with an NdeI site: CATATGGCTTCTCCTCGGAGTCCA, and a reverse primer: CCATCAACTCAATAACATAATAGACAA. The DNA sequence for the mutated form of *Adrb1* (lacking all 4 seed sites of miR-19b) was purchased from Epoch Life Science. 3' UTR fragments were then ligated into pGem-T easy vector (Promega) according to the manufacturer's guidelines, and further subcloned into a single NotI site at the 3' end of luciferase in the *Psicheck2* reporter plasmid (Promega). Cloning orientation was verified by diagnostic cuts and sequencing.

Transfections and luciferase assay

HEK293T and HT-22 cells were grown on poly-L-lysine-coated 48-well plates to a 70–85% confluence and transfected using polyethylenimine with the following plasmids: 5 ng of *Psicheck2*-3' UTR plasmid and 215 ng of EGFP overexpressing vector for either a specific miRNA, or an empty EGFP plasmid. Twenty-four hours following transfection, cells were lysed and luciferase reporter activity was assayed as previously described (Chen et al., 2005). Renilla luciferase values were normalized by control luciferase levels (transcribed from the same vector but not affected by the 3' UTR tested) and averaged across eight well repetitions per condition.

miRNA RT-qPCR expression analysis

Quantitative miRNA expression was acquired and analyzed using a step one thermocycler (Applied Biosystems), using primers that were designed for specific miRNA. RNA samples were assessed using miScript Reverse transcription kit and SYBRGreen PCR kit (Qiagen) according to the manufacturer's guidelines. U6 snRNA was used as internal control. Gene expression was obtained using the High Capacity kit and cybr green PCR master mix (Applied Biosystems). The real time PCR primers for *Adrb1* were as follows: forward: TCATGGCCTTCGTGTACTCTG; and reverse: TTACCTGTTTTGGCCCTCG.

Western blot analysis

Frozen brain samples were homogenized in RIPA buffer supplemented with proteinase inhibitors (Sigma-Aldrich) and were incubated on ice for 10 min. After 10 min centrifugation, the supernatant was transferred to a new tube and sample buffer was added to the sample, which was then boiled for 5 min and placed on ice. The samples were separated in a 10% PAGE. Transfer was performed using an assembly of nitrocellulose membrane and Whatman paper. The transfer was performed at 100 V, 350 mAmp for 1 h. After washes with PBST (PBS + 20% Tween 20)

membrane was blocked with 10% milk. The first antibody was added (rabbit anti-Adrb1, Santa Cruz Biotechnology, sc-568; mouse anti-GAPDH, Abcam, ab8245) to PBST and placed on constant shaking at 4°C overnight. The second antibody (anti-rabbit HRP, anti-mouse HRP) was added in 10% milk for 1 h. Each step was separated by additional washes with PBST. Finally, ECL was added to the membrane, which was then exposed to film.

Design, construction, and validation of miR-19 lentiviruses

miR-19b overexpression vector was cloned as follows: the enhanced form of human synapsin I promoter (Hioki et al., 2007) was PCR amplified (forward primer: tttttatcgatctcgagtagtattaatagtaac; reverse primer: tttttaccggggcgcgcccccgcagcgcagatggt) from pENTR1A-E/SYN-GFP-WRPE1 (kindly provided by Dr Takeshi Kaneko, Department of Morphological Brain Science, Graduate School of Medicine, Kyoto University, Kyoto, Japan) and inserted between ClaI and AgeI restriction sites to replace the CMV promoter in pCSC-SP-PW-GFP (kindly provided by Dr Inder Verma, The Salk Institute for Biological Studies, La Jolla, CA). Following the Synapsin promoter the precursor for miR-19b was inserted as follows: GTCCTGTATTGAGCACTGGTCTATGGTTAGTTTTGCAGGTTTGCATCCAGCTGTATAATATCTGCTGTGCA AATCCATGCAAACTGACTGTG. The H1-miR-19b KD and its control scramble sequence were obtained from GeneCopoeia and subcloned into the NheI site of the p156RRL-CMV-GFP viral plasmid. High titer lentiviruses were produced as described previously (Tiscornia et al., 2006). Briefly, recombinant lentiviruses were produced by transient transfection in HEK293T cells. Infectious particles were harvested at 48 and 72 h post-transfection, filtered through 0.45- μ m-pore cellulose acetate filters, concentrated by ultracentrifugation, redissolved in sterile HBSS, aliquot, and stored at -80°C .

Stereotactic intracranial injections

A computer-guided stereotaxic instrument and a motorized nanoinjector (Angle Two Stereotaxic Instrument, myNeuroLab) were used as previously described (Elliott et al., 2010; Kuperman et al., 2010; Regev et al., 2012). Mice were anesthetized using 1.5% isoflurane and 1 μ l of the lentiviral preparation was delivered to each basolateral amygdala using a Hamilton syringe connected to a motorized nanoinjector system at a rate of 0.2 μ l per min (coordinates relative to bregma: AP = -1.58 mm, L = ± 3.3 mm, H = -4.6 mm). Mice were subjected to behavioral studies following a 2 week recovery period.

Behavioral assessments

All behavioral assessments were performed during the dark active phase of the mice following habituation to the test room for 2 h before each test. Behavioral tests were conducted as previously described (Haramati et al., 2011; Lebow et al., 2012) in the following order, from the least stressful procedure to the most and ending with homecage locomotor testing: Open-field, dark/light transfer test, elevated plus maze, fear conditioning, and homecage locomotion.

Open-field test. The open-field test was performed in a $50 \times 50 \times 22$ cm white box, lit to 120 lux. The mice were placed in the box for 10 min. Locomotion in the box was quantified using a video tracking system (VideoMot2; TSE Systems).

Dark/light transfer test. The dark/light transfer test apparatus consists of a polyvinyl chloride box divided into a black dark compartment ($14 \times 27 \times 26$ cm) and a connected white 1200 lux illuminated light compartment ($30 \times 27 \times 26$ cm). During the 5 min test, time spent in the light compartment, distance traveled in light, and number of light/dark transitions were quantified with a video tracking system (VideoMot2; TSE Systems).

Elevated plus-maze. This apparatus in this test is designed as a plus sign and contains two barrier walls and two open arms. During the 5 min test, which is performed in relative darkness (6 lux), number of entries, distance traveled and the time spent in the open arms is automatically scored using a video tracking system (VideoMot2, TSE Systems).

Fear conditioning. A computer-controlled fear-conditioning system (TSE Systems) monitored the procedure while measuring freezing behavior (defined as lack of movement except for respiration). On the first day, mice were habituated for 5 min to the fear conditioning chamber, a

clear Plexiglas cage ($21 \text{ cm} \times 20 \text{ cm} \times 36 \text{ cm}$) with a stainless steel floor grid within a constantly illuminated (250 lux) fear conditioning housing. Conditioning took place on day 2 in one 5 min training session. Mice initially explored the context for 2 min. Thereafter, two pairings of a coterminating tone [conditioned stimulus (CS): 30 s, 3000 Hz, pulsed 10 Hz, 80 dB (A)] and shock [unconditioned stimulus (US): 0.7 mA, 2 s, constant current] with a fixed intertrial interval (ITI) of 60 s. The US was delivered through the metal grid floor. Mice were removed from this chamber 1 min after the last CS–US pairing. The chamber was cleaned thoroughly with 10% ethanol before each session. The ventilating fan of the conditioning box housing provided a constant auditory background noise [white noise, 62 dB(A)]. Context-dependent memory was tested 24 h after the conditioning by re-exposure to the conditioning box for 5 min without any stimuli. The tone-dependent cued-memory test was performed 1 d after the contextual memory test in a novel context: the walls and floor of the box were opaque black Plexiglas (dimensions were similar to the conditioning box), and the apparatus house lights and ventilating fan were turned off. Behavior was monitored for 2 min without any stimulus before the CS (tone) presentations; thereafter two CSs were presented, separated by a fixed 1 min ITI. Mice were removed from this box 1 min after the last CS.

Homecage locomotion. Homecage locomotion was assessed using the InfraMot system (TSE Systems). Mice were housed individually for 72 h, in which the first 24 h were considered habituation to the individual housing conditions. Measurements of general locomotion consisted of two light and two dark cycles in the last 48 h collected at 10 min intervals.

Microdissections, sample preparation, and HPLC-ED analysis of NE and MHPG concentrations

Microdissections were performed as previously described (Palkovits and Brownstein, 1988; Evans et al., 2008; Neufeld-Cohen et al., 2010). Briefly, coronal brain sections (300 μ m) were taken using a Leica CM1950 cryostat (North Central Instruments), mounted onto glass slides, and microdissected on a cold plate at -10°C under a stereomicroscope using microdissection needles with varying inner diameters. Each microdissected brain structure for each subject was put into separate individual tubes each containing 100 μ l of acetate buffer (3.0 g/L sodium acetate, 4.3 ml/L glacial acetic acid, pH adjusted to 5.0), which were rapidly frozen on dry ice and stored at -80°C . Next, samples were thawed and then centrifuged at 4°C and 13,000 rpm for 3 min. The supernatant was aspirated and 50 μ l of the supernatant was used for detection of noradrenaline (NE) and 3-methoxy-4-hydroxyphenylglycol (MHPG) using high performance liquid chromatography with electrochemical detection (HPLC-ED), which has been previously described (Heal et al., 1989; Evans et al., 2008), but with slight modifications. The pellet was reconstituted with 175 μ l of 0.2 M NaOH for later assay of protein content. Samples were placed in an ESA model 542 autosampler to automatically inject the samples into the HPLC system. The HPLC system also consisted of an ESA Model 582 Solvent Delivery Module to pump the mobile phase (0.1 M sodium acetate/citric acid buffer dissolved in HPLC grade H_2O , pH 4.40, adjusted with semiconductor grade NaOH, containing 8% HPLC grade methanol (v/v), and 4.6 mM octanesulphonic acid) through the chromatographic system. The stationary phase, where chromatographic separation occurred, consisted of an integrated precolumn/column system [Ultrasphere 5 ODS (C18) precolumn (45×4.6 mm)/Ultrasphere 5 ODS (C18) column (250×4.6 mm); MAC-MOD Analytical] maintained at RT. Electrochemical detection was accomplished using an ESA Model 5200A Coulochem II detector with dual potentiostats connected to an ESA 5021 Conditioning Cell with the electrode potential set at 0 mV and an ESA 5014B Microdialysis Cell with the channel 1 and channel 2 electrode potentials set at -200 and 400 mV, respectively. For each run, the average peak heights of known concentrations of NE and MHPG were determined manually using chromatography analysis software (EZChrom Elite for Windows, v2.8, Agilent Technologies) and used to calculate the concentration of the unknown samples. Tissue concentrations of NE and MHPG were standardized to the amount of protein in each microdissected structure.

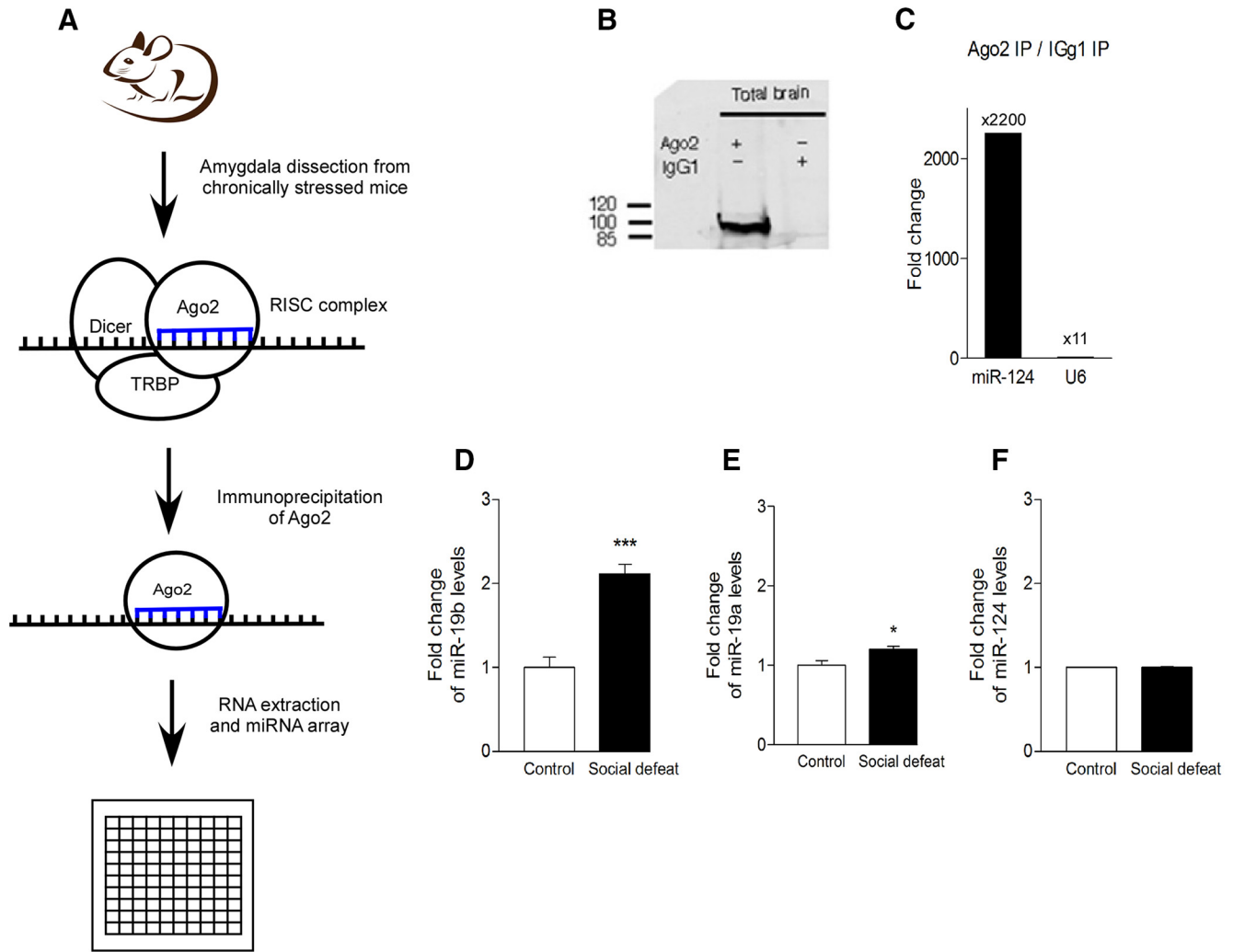


Figure 1. miR-19b is upregulated in amygdala-Ago2 extract following chronic stress. **A**, Ago2-RNA IP was performed and RNA extracted from the Ago2 complex was analyzed on an Affymetrix array. **B**, **C**, The validity of the Ago2-RNA IP was confirmed using Western blot (**B**) and RT-PCR (**C**). **D–F**, The Ago2 IP revealed a robust elevation of miR-19b (**D**) a slight elevation of miR-19a (**E**) and no change in miR-124 levels (**F**).

Statistics

The Student’s *t* test was used to assess significance. For behavioral results of lentiviral-injected mice, Student’s *t* test was used between control and miR-19b overexpression (OE) or knockdown (KD)-injected mice. Factor analysis was performed using SPSS software to determine which scores should be given to which tests. For the miRNA array results a *q* value correction was performed.

Results

MiR-19b associates with Ago2 in the amygdala following chronic stress

To identify miRs recruited to the RISC complex specifically during chronic stress we immunoprecipitated (Rodriguez-Gonzalez et al., 2008) Ago2 and its associated RNAs from the amygdala of chronically stressed mice (Fig. 1A). To validate that the immunoprecipitation indeed resulted in RNA bound to Ago2, the protein soluble fraction from total brain tissue was extracted, and the Ago2 protein-RNA complex was precipitated using anti-Ago2-coated beads. As a control we used nonspecific IgG1-coated beads (Fig. 1B). The RNA fraction was extracted from the Ago2 immunoprecipitated material by conventional methods. To verify that miRNAs were indeed precipitated with Ago2, we performed real time PCR with primers specific to miR-124 or U6. MiR-124 is

highly abundant in the brain and indeed its levels were found to be 2200-fold enriched in the Ago2 IP fraction relative to the control (Fig. 1C). The levels of U6, which is not expected to be associated with Ago2 in the RISC complex were only elevated 11-fold in the Ago2 IP (Fig. 1C) demonstrating the specificity of the Ago2 IP. To identify the entire repertoire of miRs associated with the RISC complex we performed a miRNA microarray (Exiqon) with the RNA fraction extracted from the amygdala Ago2 IP of the chronically stressed or control mice.

Analysis of the miRs that were specifically associated with Ago2 in the brains of chronically stressed mice relative to the control group identified miR-19b as the miR that showed a maximal elevation (~2-fold, $F_{(1,9)} = 41.14, p < 0.001$; Fig. 1D). This increase was specific to miR-19b, as the levels of miR-19a were only slightly elevated (~1.2-fold, $F_{(1,9)} = 9.043, p < 0.05$; Fig. 1E) and miR-124 was not significantly increased following a similar treatment (Fig. 1F). An additional group of miRs that were elevated 1.5-fold included ~30 miRs of 757 in total (data not shown).

miR-19b regulates ADRB1 levels

Bioinformatic analysis revealed ADRB1 as a possible mRNA target of miR-19b. This receptor has four seed matches for miR-19b,

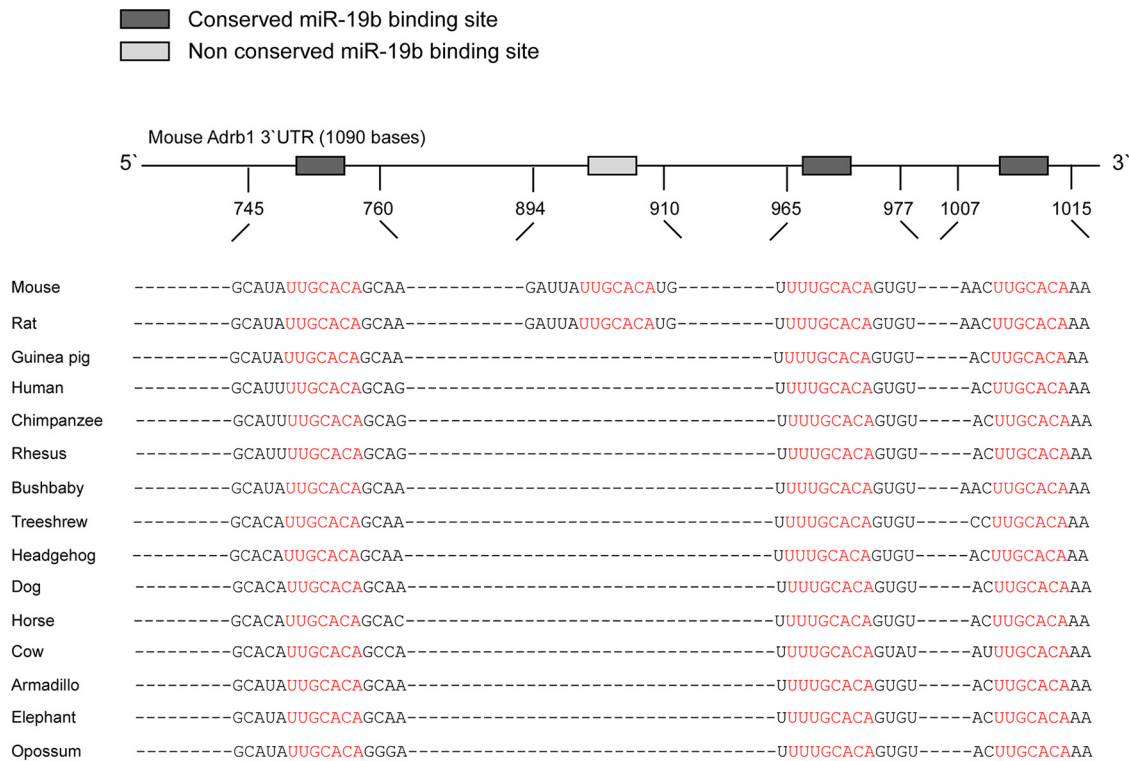


Figure 2. Schematic illustration of the 3' UTR of Adrb1 transcripts showing four putative seed matches for miR-19b and seed conservation in multiple organisms.

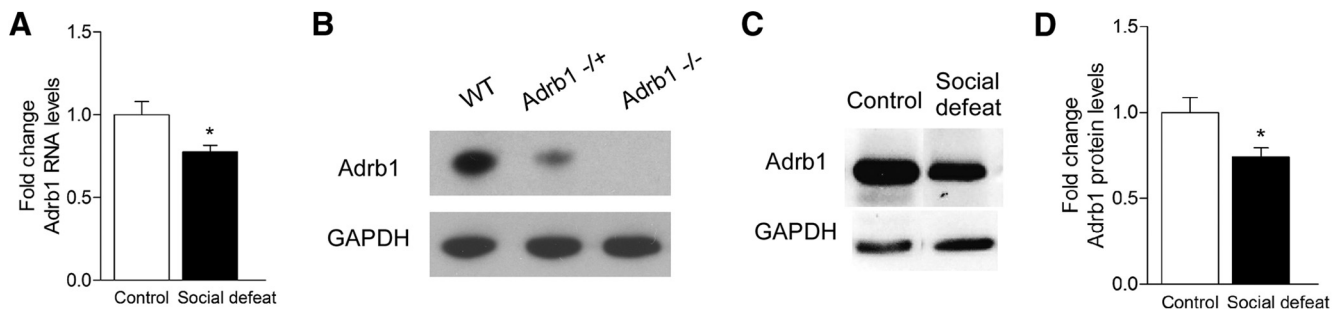


Figure 3. Amygdala Adrb1 protein levels are reduced in mice subjected to chronic social defeat paradigm. **A**, Real-time PCR analysis of Adrb1 RNA levels in the BLA of mice subjected to chronic social defeat. **B**, WB analysis of Adrb1 levels in WT, Adrb1[±] and Adrb1^{-/-} mice. **C**, **D**, WB analysis of Adrb1 protein in the BLA of mice subjected to chronic social defeat paradigm. Representative blot (**C**) and quantification of four repetitions (**D**) are presented.

three of these are conserved and one is present only in mouse and rat (Fig. 2).

Adrenergic receptors in the BLA have been reported to play a role in memory consolidation. Enhanced noradrenaline release in the BLA promotes memory consolidation. To examine whether the levels of Adrb1 are altered during social defeat in the BLA, brain tissue punches from the BLA were collected from chronically stressed and control mice and the mRNA and protein levels of Adrb1 were analyzed using real time PCR and Western blot analysis, respectively (Fig. 3). Real-time PCR analysis showed a decrease of ~25% (Fig. 3A) in the Adrb1 transcript levels of the stressed mice ($n = 5$, $F_{(1,9)} = 6.393$, $p < 0.05$). To confirm the observed changes of Adrb1 transcript levels in the protein levels, we first validated the specificity of the anti-Adrb1 antibody. Isolated BLA tissue obtained from WT, Adrb1[±] and Adrb1^{-/-} mice was analyzed using Western blot. Figure 3B clearly shows that Adrb1 is not detected in the Adrb1-deficient mice, supporting the specificity of the anti-Adrb1 antibody used. Western blot analysis showed that the levels of Adrb1 were de-

creased by ~30% (Fig. 3C,D) in the BLA of chronically stressed mice relative to control ($n = 4$, $F_{(1,9)} = 6.394$, $p < 0.05$).

To address a functional link between miR-19b and Adrb1, we expressed the luciferase gene fused to wild-type Adrb1 3' UTR or mutated Adrb1 3' UTR lacking all four seed sites (Fig. 4A) in neuronal hippocampal cell line (HT-22) and compared the luciferase activity produced by the two constructs. The HT-22 neuronal cells express very low endogenous levels of miR-19, and consequently no difference was detected between the two constructs carrying wild-type or mutated luciferase-Adrb1 3' UTR sequence (Fig. 4B). However, cotransfection of the HT-22 with miR-19b expression construct, led to reduction of ~50% ($n = 8$, $F_{(1,15)} = 69.19$, $p < 0.001$) in the luciferase activity of the luciferase-Adrb1 3' UTR relative to the control (Fig. 4C). The two luciferase-Adrb1 3' UTRs were also transfected into HEK293T cells, which express endogenously high levels of miR-19b. This resulted in a relative reduction of 50% ($n = 8$, $F_{(1,15)} = 121.348$, $p < 0.001$) in the luciferase activity in an Adrb1 3' UTR-dependent manner (Fig. 4D). Together, these experiments dem-

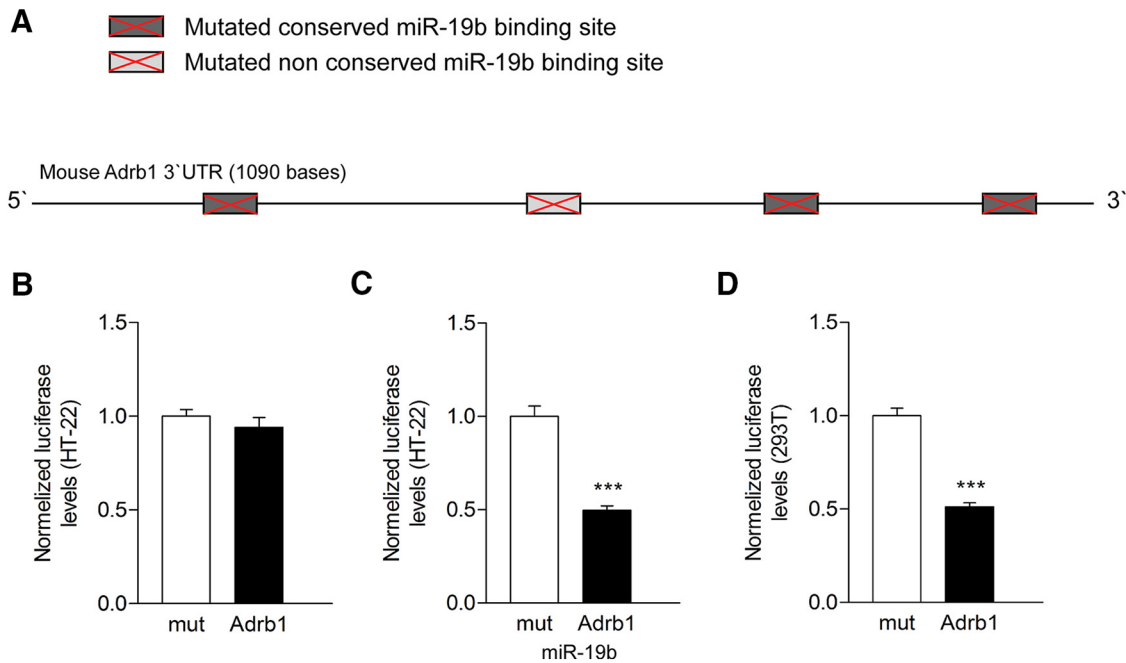


Figure 4. miR-19b regulates Adrb1 levels *in vitro* by binding to its 3' UTR. **A**, Schematic illustration of the mutated form of the Adrb1 3' UTR, lacking the four seed matches for miR-19b. **B–D**, *In vitro* luciferase assay assessing the effect of miR-19b on the 3' UTR of Adrb1 transcript. Luciferase assay in HT-22 cells that express low endogenous levels of miR-19b, without the presence of miR-19b (**B**) or in the presence of exogenous miR-19b (**C**). Luciferase assay in HEK293T cells that express high endogenous levels of miR-19b (**D**).

onstrate that miR-19b in culture conditions is capable of efficiently regulating Adrb1 through its 3' UTR.

BLA-specific miR-19b overexpression or knockdown affects auditory fear memory but not anxiety-like behavior

To examine the behavioral effects of miR-19b overexpression in the BLA of adult mice, a lentiviral vector expressing miR-19b under the neural-specific promoter synapsin was designed, constructed and produced (Fig. 5A). The precursor of miR-19b was cloned into a lentiviral construct and GFP sequences were fused at the 3'-end of the miR. A similar vector lacking the miR-19b sequences was used as a control (Fig. 5A). Viral infection of the two constructs into the hypothalamic mouse cell line N42 followed by real time PCR analysis indicated that the RNA levels of miR-19b were elevated ~6-fold ($n = 3$, $F_{(1,5)} = 148.909$, $p < 0.001$) in the infected cells (Fig. 5B), indicating an efficient production of the miR by the viral vector. A lentiviral vector expressing miR-19b knockdown or scramble sequences under the H1 promoter was designed, constructed, and produced (Fig. 5C). Infection of HEK293T cells with the knockdown viruses resulted in ~2-fold decrease in endogenous miR-19b levels relative to scrambled control ($n = 3$, $F_{(1,5)} = 13.435$, $p < 0.05$; Fig. 5D), indicating an efficient inhibition of the miR by the viral infection.

To analyze the behavioral effect of BLA-miR-19b overexpression or knockdown in adult mice, the viruses were injected bilaterally into the BLA of 10-week old C57BL/6J mice ($n = 10$ in each group; Fig. 5E–H). Following 3 weeks of recovery, the mice were subjected to the fear-conditioning test, which assess learning and memory performance following an aversive stimulus. According to previous reports (Qu et al., 2008), decreased levels of Adrb1 are associated with reduce freezing time in the fear-conditioning procedure. Based on our analysis above it was predicted that overexpression or knockdown of miR-19b in the BLA would lead to a decrease or increase of Adrb1 levels, respectively. In the first

day of the experiment (habituation) no differences were observed in the basal freezing levels between the different groups (Fig. 6A,E). Following a day of conditioning (Fig. 6B,F), in which the mice received two electrical shocks paired with a conditioned stimulus (tone), a “context test” (Fig. 6C,G) and a “cue test” (Fig. 6D,H) were performed. Freezing behavior during the context test, which are primarily associated with the hippocampus (Kim and Fanselow, 1992), was not significantly different between the overexpressing or knockdown groups of mice and their respective controls (Fig. 6C,G). However, in the cue test, we found that mice overexpressing miR-19b froze 40% less ($n = 10$, $F_{(1,19)} = 6.234$, $p < 0.05$) than the control group (Fig. 6D), suggesting a reduced memory consolidation. Intriguingly, mice injected with miR-19b knockdown viruses froze 60% more ($n = 10$, $F_{(1,17)} = 4.922$, $p < 0.05$) than the control scrambled injected group (Fig. 6H).

The overexpressing and knockdown groups of mice and their respective controls were also subjected to anxiety-like behavior assessments, including the open-field, dark/light transfer and the elevated plus maze tests. In the open-field procedure no changes were observed in time the mice spent in the center of the arena (Fig. 7A,F). Similarly, in the dark/light transfer tests, no significant differences between the groups were detected in the time spent in the light compartment (Fig. 7B,G). In addition, no significant differences between the groups were observed in the time spent in the open arms of the elevated plus-maze test (Fig. 7C,H), although the miR-19b knockdown group showed a strong tendency ($n = 10$, $F_{(1,19)} = 4.243$, $p = 0.054$) for increased time in the open arms (Fig. 7H). No significant differences were observed in the homecage locomotor activity of the tested groups of mice (Fig. 7D,E,I,J). Together, the behavioral experiments support a specific role for miR-19b in regulating auditory fear memory, which is potentially mediated by the influence of miR-19b on Adrb1.

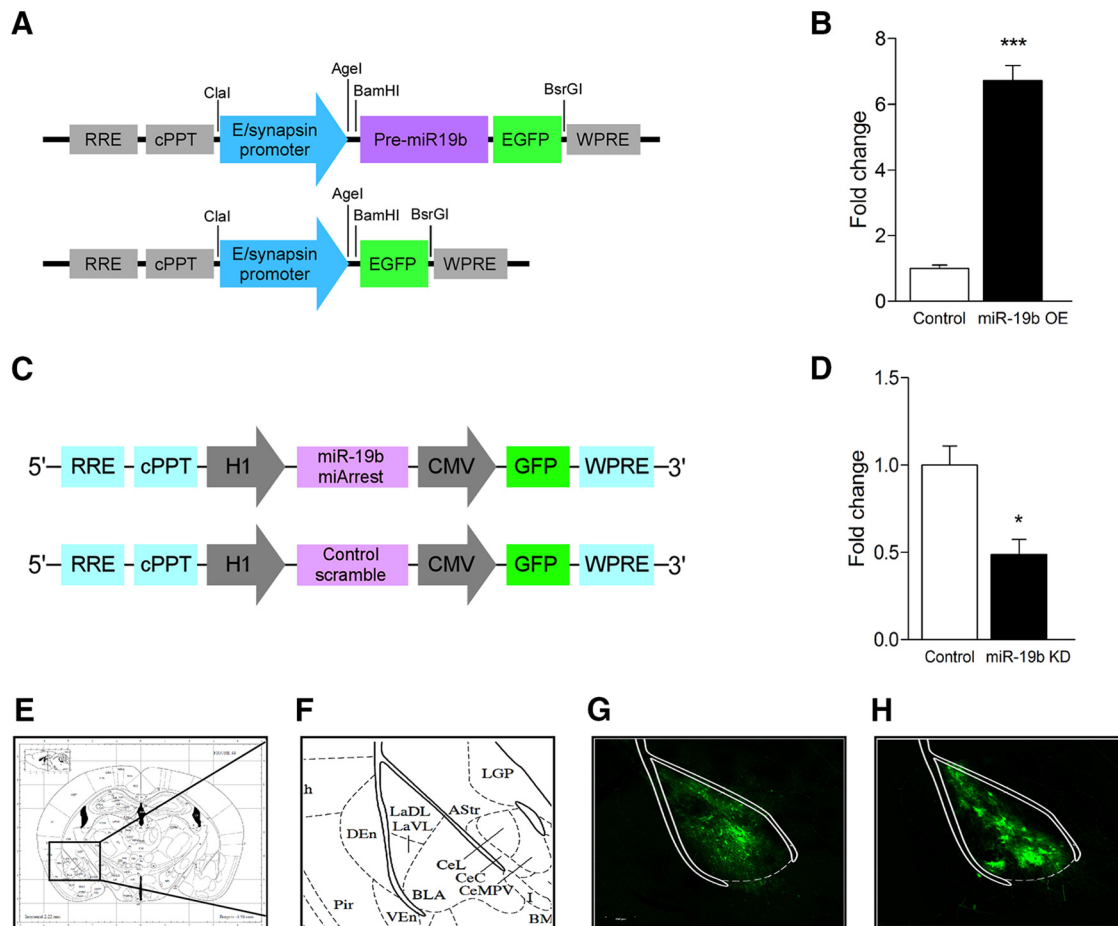


Figure 5. Establishment and validation of miR-19b overexpressing and knockdown lentiviruses. **A**, Schematic illustration of the Syn-miR-19b OE and control lentiviral constructs. **B**, Infection of N42 cells with the Syn-miR-19b OE virus showed an elevation of ~6-fold in miR-19b levels. **C**, Schematic illustration of the miR-19b knockdown (KD) and control lentiviral constructs. **D**, Infection of HEK293T cells with the miR-19b KD virus showed a decrease of ~2-fold in miR-19b levels. **E**, Schematic representation of site of delivery adopted from Paxinos and Franklin digital mouse brain atlas. **F**, Enlargement of the basolateral amygdala region corresponding to the injection site. **G**, **H**, Representative microscope image of virally infected BLA at 10 weeks following miR-19b OE (**G**) or miR-19b KD (**H**) virus injection.

Downstream effects of BLA-specific miR-19b regulation on limbic circuits

The effects of noradrenaline-induced BLA activation on memory consolidation are thought to be mediated by downstream signaling to limbic forebrain structures including the hippocampus and medial prefrontal cortex (Schwabe et al., 2012). Therefore, we predicted that BLA-specific modulation of miR-19b might alter signaling in downstream limbic structures. Noradrenaline acts via β adrenergic receptors within the BLA to enhance excitatory neurotransmission both *in vivo* (Buffalari and Grace, 2007) and *in vitro* (Gean et al., 1992; Huang et al., 1996; Ferry et al., 1997). Because noradrenaline acts via β adrenergic receptors within the BLA to activate glutamatergic projection neurons, and glutamate acts on presynaptic noradrenergic terminals to induce noradrenaline release in a number of limbic forebrain structures, including the prefrontal cortex (Russell and Wiggins, 2000) and hippocampus (Howells and Russell, 2008; Dazzi et al., 2011), we predicted that selective overexpression of miR-19b, by downregulating *Adrb1* and noradrenaline-mediated activation of glutamatergic projection neurons, would decrease noradrenergic signaling in specific downstream targets of the BLA. Complementarily, knockdown of miR-19b in the BLA may possibly result in an opposite affect by increasing *Adrb1* levels.

Following the behavioral and locomotor activity tests, miR-19b overexpressing or knockdown injected mice were sacrificed and noradrenaline and its degradation metabolite, MHPG, were measured in specific brain nuclei (Fig. 8A–C). A treatment \times region interaction for the MHPG:NE ratio ($F_{(4,56,82.0)} = 3.38$; $p = 0.01$; $\epsilon = 0.285$) was observed. The BLA-specific miR-19b overexpression decreased the MHPG:NE ratio in the dorsal hippocampus (CA1d region) by ~20% ($F_{(1,17)} = 5.213$, $p < 0.05$) as expected (Fig. 8D), suggesting that BLA-specific miR-19b overexpression not only influences noradrenergic signaling within the BLA, but also in a distributed system controlling memory consolidation (Roosendaal, 2000). Interestingly, the miR-19b knockdown group showed increase of ~25% in the MHPG:NE ratio in the same brain region (CA1; $F_{(1,21)} = 5.916$, $p < 0.05$; Fig. 8G). The MHPG:NE ratios in the prelimbic (PrL) and infralimbic (IL) structures were unchanged (Fig. 8E, F, H, I). There was no treatment or treatment \times region interactions for either the MHPG or NE concentrations alone.

Discussion

Adrb1 was previously demonstrated to play a pivotal role in regulating memory consolidation and stabilization (Roosendaal et al., 2004, 2006a) and anxiety-like behavior (Rudoy and Van Bockstaele, 2007; Fu et al., 2008). Therefore, modulating *Adrb1*

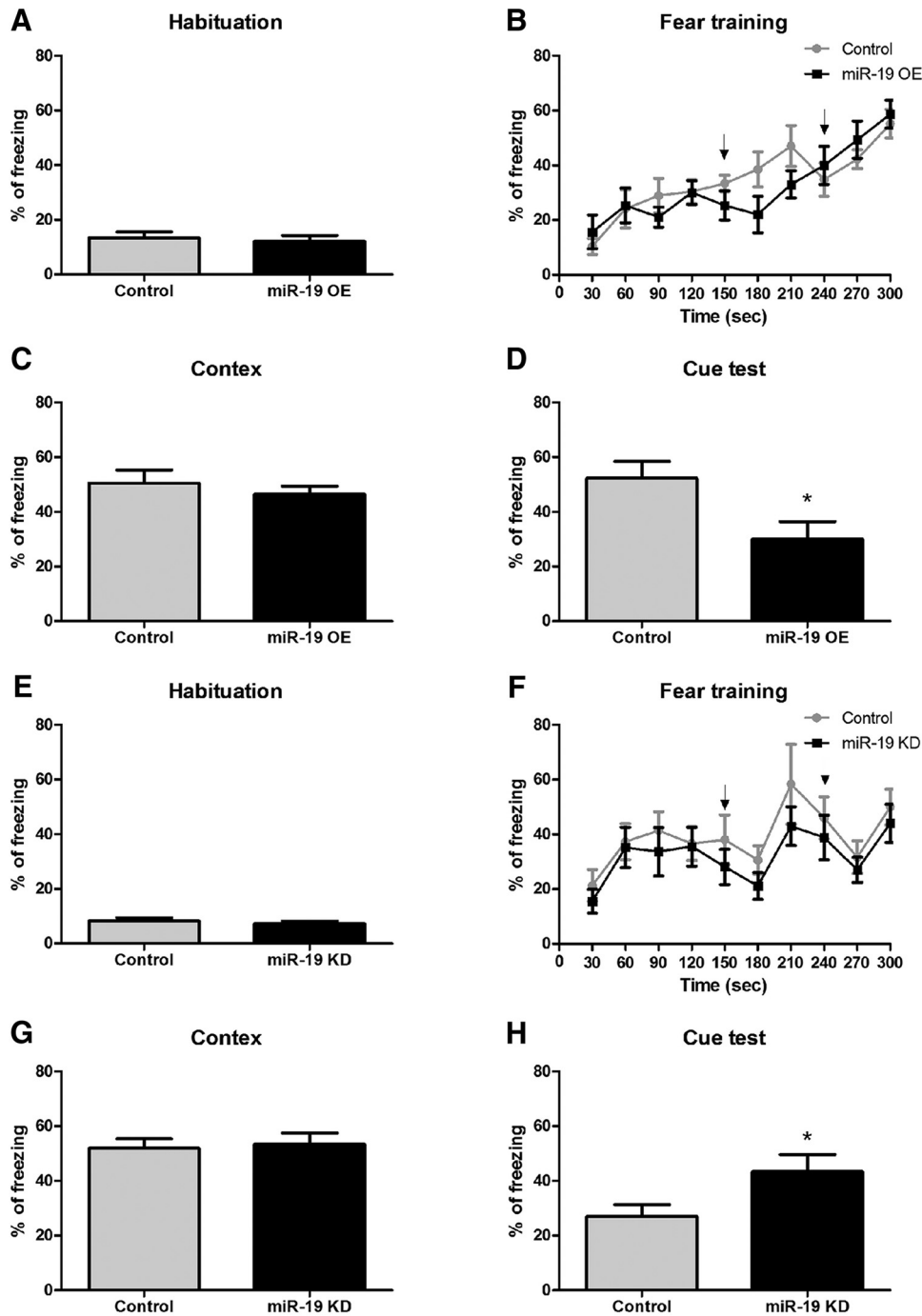


Figure 6. BLA-specific miR-19b overexpressing or knockdown mice showed changes in memory consolidation. *A–H*, Percentage of freezing in the fear conditioning paradigm ($n = 10$). Although no significant changes were observed in the percentage of freezing in the habituation (*A, E*), fear training (*B, F*), or context test (*C, G*), the BLA-specific miR-19b OE mice showed significantly reduced percentage freezing in the Cue test (*D*), whereas the BLA-specific miR-19b knockdown (KD) mice showed significantly elevated percentage freezing in the Cue test (*H*).

levels and/or activity was proposed to be important for the regulation of stress-induced behavioral responses (Fortaleza et al., 2012). Here we identify miR-19b as an important novel modulator of *Adrb1* in the BLA. MiR-19b levels were found to be significantly elevated in the RISC RNA-Ago2 complex in the amygdala region following the application of chronic social defeat (CSD) in mice, concomitant with a significant reduction in the transcript and protein levels of its putative target, *Adrb1*. These findings suggested a role for miR-19b in regulating amygdalar transcripts following chronic stress, and revealed *Adrb1* as a key target and mediator of this process. Indeed our results demonstrate *Adrb1*

as a major target of miR-19b regulation. MiR-19b expression reduced the levels of a luciferase reporter fused to the 3' UTR of *Adrb1* in a manner that was dependent on the presence of four miR-19b seeds in its 3' UTR. Furthermore, changing the levels of BLA-miR-19b in adult mice using site-specific viral approach resulted in changes in freezing duration in the cue test of the fear-conditioning procedure. Whereas higher levels of miR-19b in the BLA resulted in reduced freezing in the cue test, lower levels of BLA-miR-19b caused an opposite effect, showing increased freezing behavior. These behavioral changes were also reflected in the opposed MHPG:NE ratio in the dorsal hippocampus suggest-

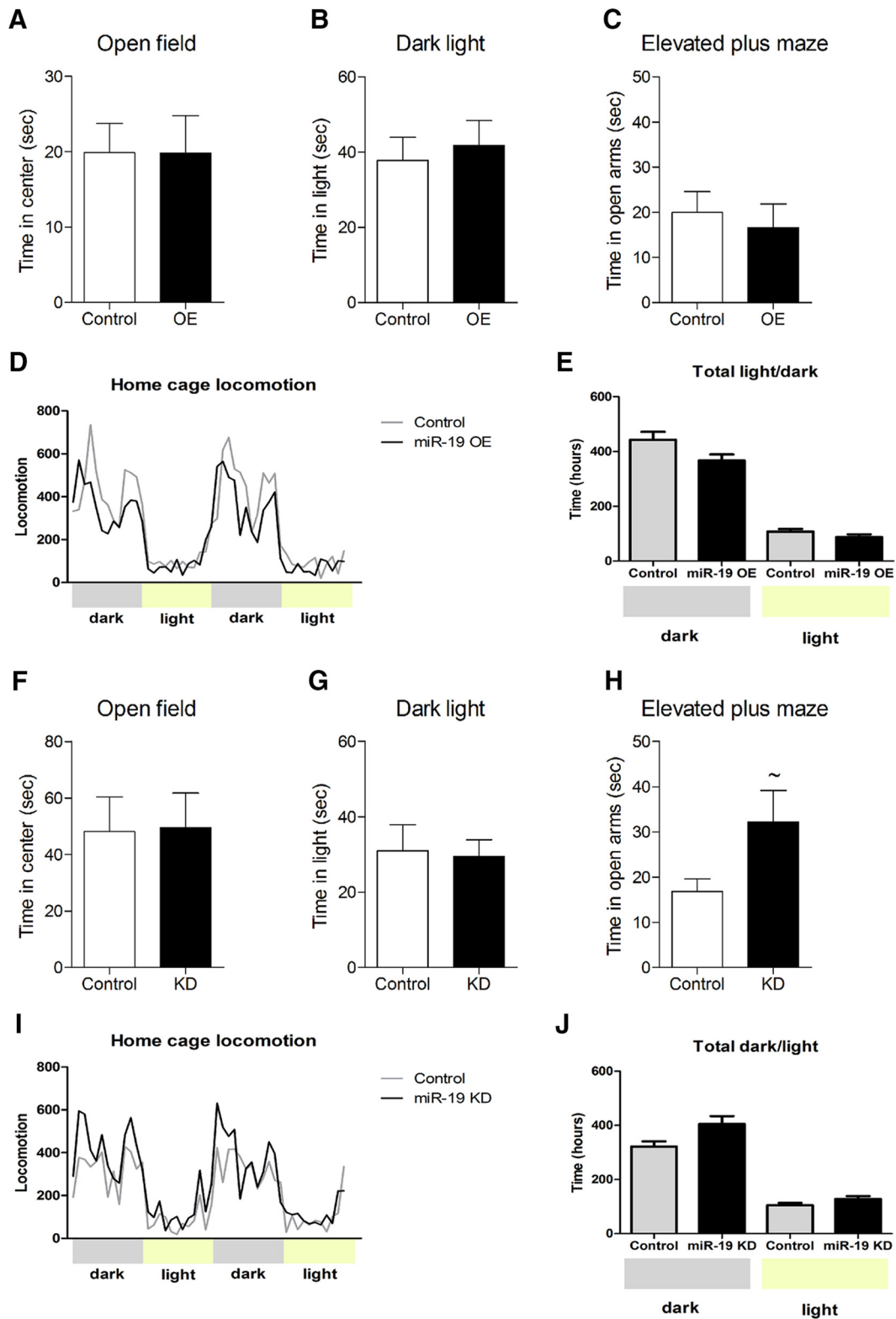


Figure 7. Intact anxiety-like behavior in mice expressing modified levels of miR-19 in the BLA. **A–C, F–H**, Anxiety-like behavior was assessed using the open-field (**A, F**), dark/light transfer (**B, G**), and elevated plus maze (**C, H**) tests. The BLA–miR-19b OE or miR-19b KD groups of mice ($n = 10$) showed no significant differences in the anxiety parameters examined. **D, E, I, J**, No significant changes were observed in the homecage locomotor activity between the BLA–miR-19b OE mice and the control group (**D, E**) and the BLA–miR-19b KD mice and their respective control (**I, J**).

ing modified noradrenergic signaling in a distributed system controlling memory consolidation. These results suggest that BLA–miR-19b levels are important for the regulation of memory stabilization in response to stressful challenge, which is mediated

primarily via modulating *Adrb1* levels. The high BLA–miR-19b levels in the Ago2 complex of chronically stressed mice reported in this study may be directly linked to the long-term regulation of the central stress response and related copying mechanisms. The

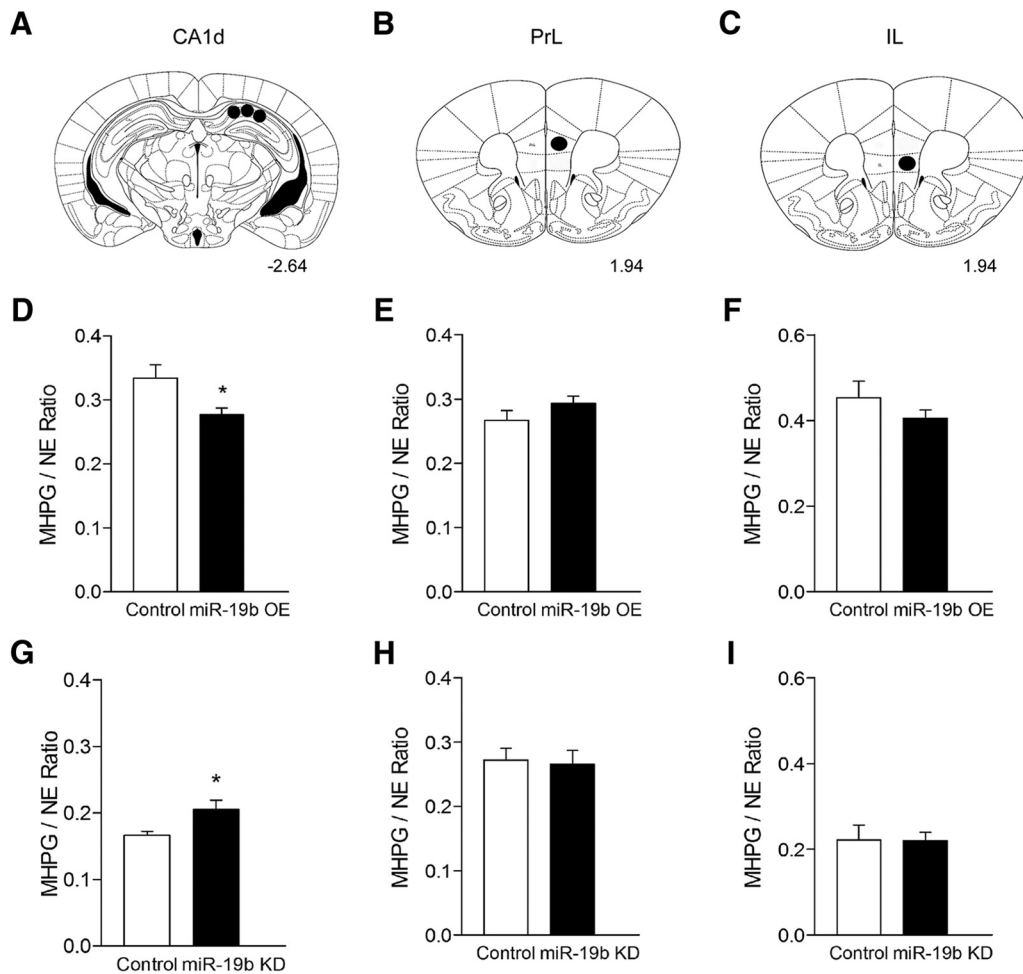


Figure 8. miR-19b OE and KD mice showed a reduction of MHPG:NE ratio in the hippocampus. **A–C**, Schematic illustration adopted from Paxinos and Franklin digital mouse brain atlas showing the nuclei analyzed in the MHPG:NE ratio measurements. Analysis of the MHPG:NE ratio in the CA1d (**D**, **G**), PrL (**E**, **H**), and IL (**F**, **I**) brain regions ($n = 10$).

regulation of *Adrb1* levels may be important for the coping of mice with stressful challenges. Low levels of *Adrb1* that mediate a decrease in memory consolidation might pan out as beneficial to accommodate stressful events. Several previously published studies have highlighted adrenergic dysregulation in PTSD human studies (Southwick et al., 1999; Strawn and Geraciotti, 2008; Krystal and Neumeister, 2009; Liberzon et al., 2014), suggesting an important role for the adrenergic system in coping with stressful events.

Social defeat mice have been shown previously to exhibit higher freezing time in the auditory cue test (Yu et al., 2011). Our current findings show that as a consequence of chronic stress, miR-19b levels are elevated in the amygdala, and when chronically overexpressed in the BLA caused lowering of the freezing time in the cue test. Although these findings may appear as contradicting, the experimental approaches are fundamentally different and the final behavioral readout could be influenced by multiple different factors. The CSD protocol is associated with multiple molecular and cellular changes that collectively translated, among others, into increase freezing time in the auditory cue test. Although miR-19b levels are upregulated following exposure to CSD protocol, the overexpression of miR-19b in naive mice are mechanistically very different from exposing the mice to CSD. Clearly, the overexpression of BLA-miR-19b does not represent all the changes occurring following CSD protocol, which will be also reflected in the behavioral and physiological phenotype of these mice. Additional possible factor that may contribute

for these differences is the expression kinetics of the endogenous miR-19b in the BLA following stressful challenge. Here, we demonstrate a significant increase in miR-19b in the BLA of mice 8 d following the termination of the CSD protocol. This upregulation could be transient, permanent, or even exhibit a more complex pattern of expression kinetics and therefore the constitutive overexpression of the virally mediated miR-19b in the BLA not necessarily mimicked the endogenous situation. Nevertheless, the demonstrated repressive effect of miR-19b on *Adrb1* levels and its downstream behavioral and cellular effects persist.

Whereas the three subtypes of the β adrenergic receptors are expressed in different tissues, i.e., beta1 is expressed by heart cardiomyocytes, beta2 is the dominant subtype distributed in the lungs (Carstairs et al., 1985), and beta3 is expressed in adipose tissue (Krief et al., 1993), all three are expressed in the CNS (Insel, 1993). *Adrb1* and *Adrb2* are both expressed in neurons of the BLA and not in astrocytes. In addition, whereas *Adrb1* is uniquely detected in the cytoplasm, *Adrb2* has been shown to be present in both the cytoplasm and nucleus (Qu et al., 2008). This may indicate that each of these receptors have a distinct mechanism of action and may be differentially regulated. *Adrb2* has been shown to be controlled *in vitro* in lung epithelial H292 cells by the miR *let-7f*. *Let-7* expression increases with age (Nishino et al., 2008) and thus its regulation of *Adrb2* in the BLA may be mediated through an age-related mechanism, although our analysis shows that *Adrb1* is regulated in the BLA by stress. Previously, it has

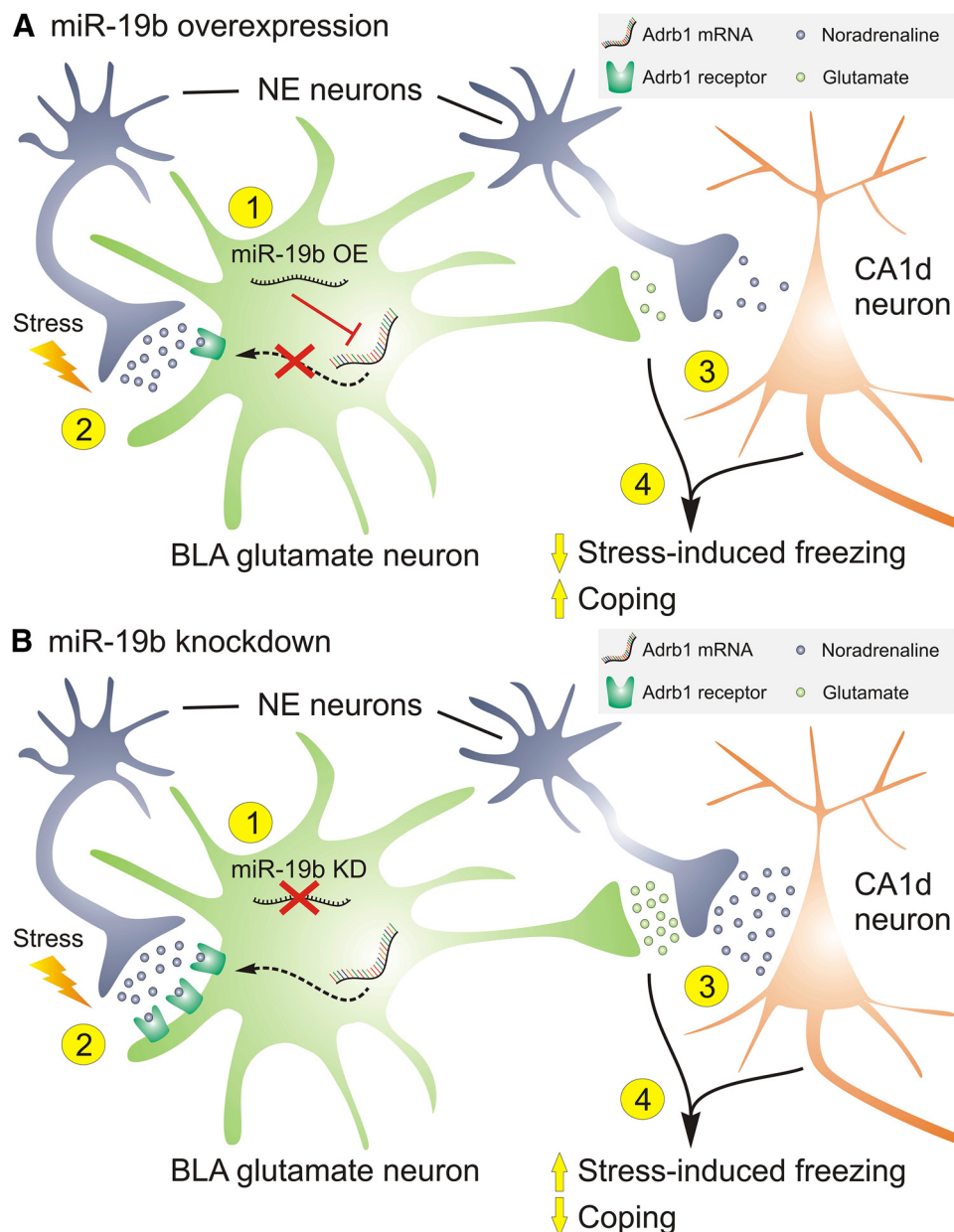


Figure 9. Putative model illustrating the effects of either miR-19b OE (**A**) or miR-19b knockdown (KD; **B**) in the BLA on NE neurotransmission in downstream limbic structures, such as CA1d and the resulting changes in coping behavior. Stress-induced activation of NE afferents (Morilak et al., 2005; Itoi and Sugimoto, 2010) results in excitation of subpopulations of BLA principle neurons through agonism of β -adrenergic receptors (Ferry and McGaugh, 2000; Buffalari and Grace, 2007); therefore, in **A**, (1) miR-19b OE in the BLA reduces Adbl1 signaling through RNA transcript degradation or translation inhibition, which leads to (2) reduced stress-induced activation of BLA glutamatergic projection neurons by NE afferents. Together with evidence that BLA glutamatergic projection neurons target the hippocampal formation (Sparta et al., 2014), including the CA1d region (Pikkarainen et al., 1999; Pitkänen et al., 2000), and these neurons are thought to be important for fear-related behavioral responses and memory processes (Davis, 1992; Herry et al., 2008; Zimmerman and Maren, 2010), as well as evidence that glutamate regulates hippocampal NE release through modulation of presynaptic NE terminals (Russell and Wiggins, 2000; Luccini et al., 2007; Howells and Russell, 2008; Dazzi et al., 2011), (3) it is probable that the decreased stress-induced activation of BLA glutamatergic projections results in diminished stress-induced presynaptic stimulation of NE release in the CA1d and consequently, and (4) reduced stress-induced freezing in cued fear conditioning and increased coping. Conversely, in **B**, (1) miR-19b KD in the BLA facilitates Adbl1 signaling resulting in (2) increased stress-induced activation of BLA glutamatergic projection neurons by NE afferents. (3) The greater stress-induced activation of BLA glutamatergic projections results in augmented stress-induced presynaptic stimulation of NE release in the CA1d and consequently, and (4) increased stress-induced freezing and decreased coping.

been shown that the blockade of Adbl1 and Adbl2 in the BLA immediately after conditioning of rats resulted in a significantly lower freezing score in the auditory test performed 24 h postconditioning, indicating that both receptors are important for the formation of auditory fear memory (Qu et al., 2008). Our data implicate that miR-19b targets specifically Adbl1 and not Adbl2, suggesting a nonredundant function for Adbl1 in the stress response. This is consistent with studies by Qu et al. (2008) dem-

onstrating that blocking either Adbl1 or Adbl2 was sufficient to cause a severe deficit in auditory fear memory. Consistent with blocking specifically neuronal but not astrocyte Adbl1, the virus used for injection in our miR-19b overexpressing experiments included a neuronal specific synapsin promoter. However, it is conceivable that the injection of miR-19b into the BLA affected other potential mRNA targets in the neurons. For example, miR-19b can potentially target the cannabinoid receptor type 1 (CB1).

CB1 has a 3' UTR of 3.8 kb with two seed matches for miR-19b (one conserved and one less conserved). CB1 plays an important role in memory consolidation in the model proposed by Hill and McEwen (2009) by releasing GABA inhibition in the BLA neurons and inducing noradrenaline release, promoting memory consolidation in the BLA.

Whereas the mice injected with miR-19b overexpressing or knockdown viruses showed a significant difference in freezing time in the cue test, no changes were observed in their freezing duration in the habituation period. Similarly, no changes were observed in their locomotor activity at the anxiety-like behavior tests or the homecage locomotor measurements. Both observations indicate that there were no motor differences between the groups and the change in freezing time resulted from a specific change in memory consolidation rather than alterations in motor function.

MiR-19b might be part of a group of miRs controlling different target genes that contribute to memory consolidation. For example, overexpression of miR-182 has been implicated recently in disruption of memory consolidation in the lateral amygdala of rats (Griggs et al., 2013). Griggs et al. (2013) showed that overexpression of miR-182 in the lateral amygdala represses expression of cortactin and Ras-related C3 botulinum toxin substrate 1 (Rho family, small GTP binding protein Rac1; Rac1) following auditory fear conditioning. The levels of miR-128b were demonstrated to be elevated in the infralimbic prefrontal cortex following fear extinction learning in mice, suggesting a role for this miR in facilitating the formation of a fear extinction memory through targeting of the plasticity-related gene, regulator of calmodulin signaling (Rcs; Lin et al., 2011). Finally, miR-132 overexpression was demonstrated to cause impaired plasticity and a functional deficit in short-term recognition memory (Scott et al., 2012).

In summary, we have identified miR-19b as an important chronic stress-induced regulator of amygdalar *Adrb1* and its subsequent behavioral functions. A putative model illustrated in Figure 9 describe the effects of either miR-19b overexpression or miR-19b knockdown in the BLA on NE neurotransmission in downstream limbic structures, such as the CA1 region of the dorsal hippocampus, and the resulted changes in coping behavior. MiR-19b overexpression in the BLA reduces *Adrb1* signaling through RNA transcript degradation or translation inhibition, which leads to reduced stress-induced activation of BLA glutamatergic projection neurons by NE afferents. The decreased stress-induced activation of BLA glutamatergic projection results in diminished stress-induced presynaptic stimulation of NE release in the CA1d and consequently reduce stress-induced freezing in the cued fear conditioning and increase coping. Conversely, miR-19b knockdown in the BLA facilitates *Adrb1* signaling resulting in increased stress-induced activation of BLA glutamatergic projection neurons by NE afferents. The greater stress-induced activation of BLA glutamatergic projections results in augmented stress-induced presynaptic stimulation of NE release in the CA1d and consequently increase stress-induced freezing and decrease coping. Spatial regulation of *Adrb1* by miRNA enables the neurons in the BLA to provide an appropriate response in stress conditions enabling the animal to accommodate its behavior to the changing conditions.

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