Systems/Circuits

Adolescent Parvalbumin Expression in the Left Orbitofrontal Cortex Shapes Sociability in Female Mice

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The adolescent social experience is essential for the maturation of the prefrontal cortex in mammalian species. However, it still needs to be determined which cortical circuits mature with such experience and how it shapes adult social behaviors in a sex-specific manner. Here, we examined social-approaching behaviors in male and female mice after postweaning social isolation (PWSI), which deprives social experience during adolescence. We found that the PWSI, particularly isolation during late adolescence, caused an abnormal increase in social approaches (hypersociability) only in female mice. We further found that the PWSI female mice showed reduced parvalbumin (PV) expression in the left orbitofrontal cortex (OFC_L). When we measured neural activity in the female OFC_L , a substantial number of neurons showed higher activity when mice sniffed other mice (social sniffing) than when they sniffed an object (object sniffing). Interestingly, the PWSI significantly reduced both the number of activated neurons and the activity level during social sniffing in female mice. Similarly, the CRISPR/Cas9-mediated knockdown of PV in the OFC_L during late adolescence enhanced sociability and reduced the social sniffing-induced activity in adult female mice via decreased excitability of PV^+ neurons and reduced synaptic inhibition in the OFC_L. Moreover, optogenetic activation of excitatory neurons or optogenetic inhibition of PV^+ neurons in the OFC_L enhanced sociability in female mice. Our data demonstrate that the adolescent social experience is critical for the maturation of PV^+ inhibitory circuits in the OFC_L; this maturation shapes female social behavior via enhancing social representation in the OFC_L.

Key words: adolescent social isolation; female; inhibition; maturation; orbitofrontal cortex; parvalbumin

Significance Statement

Adolescent social isolation often changes adult social behaviors in mammals. Yet, we do not fully understand the sex-specific effects of social isolation and the brain areas and circuits that mediate such changes. Here, we found that adolescent social isolation causes three abnormal phenotypes in female but not male mice: hypersociability, decreased PV^+ neurons in the left orbitofrontal cortex (OFC_L), and decreased socially evoked activity in the OFC_L. Moreover, parvalbumin (PV) deletion in the OFC_L *in vivo* caused the same phenotypes in female mice by increasing excitation compared with inhibition within the OFC_L. Our data suggest that adolescent social experience is required for PV maturation in the OFC_L, which is critical for evoking OFC_L activity that shapes social behaviors in female mice.

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Introduction

One of the hallmarks of postnatal maturation in the mammalian cortex is the maturation of GABAergic inhibitory circuits (Luhmann and Prince, 1991). Maturation is accompanied by an increase in the expression of both the parvalbumin (PV) in a subset of GABAergic neurons and the perineuronal nets (PNNs) surrounding the PV⁺ neurons (Celio and Blumcke, 1994; Huang et al., 1999). PV is a calcium-binding protein and a well known genetic marker for fast-spiking GABAergic neurons in the cortex (Rudy et al., 2011). PV gene expression is upregulated in the sensory cortex in an activity-dependent manner (Sugiyama et al., 2008; Krishnan et al., 2017) after mammals are born. Like PV, PNNs also increase during cortical maturation and, in the adult cortex, surround most of the PV⁺ neurons (Ueno

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et al., 2017, 2018). Along with the increase of PNNs in cortical circuits, other molecular changes reduce net plasticity and facilitate the maturation of the cortical network (Beurdeley et al., 2012). If mice are deprived of visual input during the critical period after eye opening, via monocular deprivation or dark rearing, PV^+ neurons are less developed, and PNNs are reduced in the primary visual cortex (V1; Sugiyama et al., 2008). This deprivation, in turn, diminishes both the visual response of V1 neurons that receive input from the deprived eye (Sugiyama et al., 2008) and inhibition in the cortex (Dityatev et al., 2007). Therefore, the postnatal sensory experience is critical for the expression of PV and PNNs in PV⁺ neurons, as these are required for the maturation of PV⁺ inhibitory circuits and, eventually, for shaping sensory responses in the cortex (Hensch, 2005).

The maturation of the cortex occurs first in the sensory area, where the process is part of sensory experience, and later in the motor-frontal and the prefrontal cortex (PFC; Hensch, 2005). The social experience is critical for the maturation of the PFC in both humans (Blakemore, 2008; Paus et al., 2008) and rodents (Makinodan et al., 2012; Bicks et al., 2020; Tan et al., 2021), particularly during juvenile and adolescent periods. Patients with autism or schizophrenia often show abnormal social cognition as well as dysfunction in the PFC (Bicks et al., 2015). In these patients, the expression of PV and PNNs is reduced in the dorsolateral PFC (Enwright et al., 2016). Thus, it is plausible that immature PV⁺ circuits in the PFC cause the psychiatric symptoms underlying abnormal social behavior in mammals. However, we still do not understand whether social experience is required for PV⁺ inhibitory circuits in these PFC regions to mature. In particular, it is largely unknown whether social experience affects the maturation of inhibitory circuits in the orbitofrontal cortex (OFC), one of the PFC subregions critical for social cognition (Adolphs, 2003; Amodio and Frith, 2006).

Mammalian social behavior is important for maintaining a conspecific society. Social interaction during adolescence is critical for shaping social and cognitive behaviors in both humans and rodents (Spear, 2000; Makinodan et al., 2012; Hinton et al., 2019; Almeida et al., 2021; Potrebic et al., 2022). Interestingly, the mammalian brain shows hemispheric asymmetry and lateralization in executing social behaviors (Marlin et al., 2015). Even in rodents, according to recent studies, the PFC functions asymmetrically in mediating social behaviors (Lee et al., 2015; Marlin et al., 2015). However, whether such asymmetry in the PFC becomes increasingly pronounced in a sex-specific manner with social experience during adolescence is still unclear. We investigated the effect of adolescent social isolation on the maturation of PV⁺ neurons in the PFC in both sexes and hemispheres. Using postweaning social isolation (PWSI) and the CRISPR/ Cas9 system, we revealed that social isolation during adolescence caused the defect in PV expression in the left OFC (OFC_L) of female mice, which led to hypersociability and reduced OFC_L activity during social contact.

Materials and Methods

Experimental model and subject details

We approved all experimental procedures from the KAIST Institutional Animal Care and Use Committee (IACUC-18–236). We maintained all mice *ad libitum* under light from 8:00 A.M. to 8:00 P.M. and dark from 8:00 P.M. to 8:00 A.M. All mice were weaned at postnatal day 21 (P21); mice were reared in double cages $(28 \times 17 \times 14 \text{ cm})$ with three to five mates for group housing or in single cages $(20 \times 10 \times 14 \text{ cm})$ alone for social isolation. We performed experiments on WT (C57BL/6J), PV-Cre

(catalog #008069, The Jackson Laboratory), and the Rosa26-floxed STOP-Cas9 knock-in mice (catalog #024857, The Jackson Laboratory).

Histology

We anesthetized mice with isoflurane and then perfused them with $1 \times$ PBS and 4% paraformaldehyde (w/v in PBS). Mouse brain samples were postfixed for 4 h at 4°C; after fixation, we incubated them in 30% sucrose solution (w/v in PBS) for 1-2 d at 4°C. Then, we embedded each brain sample in O.C.T. medium (catalog #4583, Tissue-Tek) and froze each at -80°C. We sectioned frozen brains using a cryostat (Leica) and collected coronal sections either at 30 μ m (for confirming injection sites) or 15 μ m (for immunostaining) thickness. To confirm fluorescently labeled injection sites, brain sections were washed three times with $1 \times PBS$ and mounted in DAPI-containing mounting medium (VECTASHIELD, catalog #H-1200, Vector Laboratories). For immunostaining, we washed the brain sections three times with $1 \times PBS$ and permeabilized them with 0.3% Triton X-100 in $1 \times$ PBS for 30 min. After the sections were incubated in a blocking solution (2% normal donkey serum in $1 \times PBS$) for 2 h at room temperature (RT), we incubated them with primary antibodies (1:500 dilution in blocking solution; rabbit anti-PV; catalog #PV25, Swant; 1:500 dilution in blocking solution; mouse anti-NeuN; catalog #MAB377, Millipore) for 2 d at 4°C. Last, we incubated them with fluorescence-tagged secondary antibodies (Alexa Fluor 594 donkey anti-rabbit IgG for PV staining; 1:500 dilution in 1× PBS; catalog #A21207, Thermo Fisher Scientific; Alexa Fluor 647 donkey anti-mouse IgG for NeuN staining; 1:500 dilution in $1 \times$ PBS; catalog #715–605-151, Jackson ImmunoResearch). For PNN staining, we treated the brain sections with the fluorescein (FITC)-labeled Wisteria floribunda agglutinin (WFA; 1:500 dilution in $1 \times$ PBS; catalog #FL-1351, Vector Laboratories) for 2 h at RT before mounting them in clear mounting medium (catalog #S3023, DAKO). We imaged fluorescent signals of mounted brain sections under a scanning microscope (Axio Scan.Z1, Zeiss).

Cell-counting analysis

When we sectioned the mouse brain samples, we collected 15 µm brain sections in every 60 µm thickness across the frontal cortex. After immunostaining the sections, we scanned the fluorescent images of the brain sections and matched the images to the mouse brain atlas (Franklin and Paxinos, 2012) to confirm the anterior-to-posterior location of the coronal brain sections. We then selected three images of sections in a row that included OFC areas (between +2.10 mm and approximately +2.34 mm anterior from the bregma; see Fig. 2C). To count the labeled neurons in the OFC, we placed a square $(0.3 \times 0.3 \text{ mm}^2)$ as a region of interest (ROI) at the center of the OFC (± 1.3 mm lateral from the bregma; depth, -2.5 mm). To quantify the number of PV⁺ and NeuN⁺ neurons, we used automatic cell counting software (ImageJ; Grishagin, 2015). After converting the fluorescent images to 8 bit grayscale images, we further transformed the grayscale images into binary images. For the binary transition, we used distinct thresholds at the grayscale for isolating the NeuN⁺ neurons (see Fig. 2; total mean intensity, $11.39 \pm 0.02\%$) and the PV^+ neurons (see Fig. 2; total mean intensity, 2.12 \pm 0.16%). We then automatically counted the white-filled circles as cells if they satisfied the following conditions: (1) the total size was $>60 \times 60$ pixels; and (2) the circularity was >0.2. We set the thresholds in each sample at the level that yielded a similar number of cells from the automatic counting compared with the manual counting using a couple of template images from each sample. There was no significant difference between the number of cells counted automatically and the number of cells counted manually in a blind manner (see Fig. 2B). To validate the knock-down efficiency of single-guided RNAs targeting the parvalbumin gene or chondroitinase ABC (chABC; see Fig. 5), we manually and blindly counted the number of PV⁺ or WFA⁺ neurons in the ROI.

Sociability test

We performed the sociability test by using the following two types of mazes: one was a modified three chamber-like maze ($50 \times 45 \times 20$ cm) where two containers were located diagonally at opposite corners; and the other was a standard three-chamber maze ($60 \times 40 \times 20$ cm), which was used in other studies (Won et al., 2012). Most of the data we plotted

in the article were collected in the modified maze. We performed the sociability test on mice at either P79 to P81 or P50. Before behavior experiments, all mice were habituated for 30-60 min in the habituation chamber. All mice were maintained in a group-housed (GH) or an isolated condition until the habituation. Mice were then moved to the sociability test maze. The sociability test consisted of the following two phases: (1) habituation in the sociability maze (empty containers, 10 min); and (2) sociability test (an object container and a social container with a stranger mouse, 10 min). We used a 2 week younger and same-sex mouse as a stranger mouse not to evoke any aggressively or sexually driven social behaviors in the tested mouse (Murugan et al., 2017). We used the software EthoVision XT 11.5 (Noldus) to automatically analyze interaction times and manually analyze the sniffing time. The sniffing time was defined as the time when the nose of the subject mouse contacted the container either empty or with a novel mouse. We analyzed mouse sociability by the following two indicators: time spent in interaction zones near the container (18 \times 18 cm) and sniffing time. We calculated the social-sniffing and social interaction indices as follows (Nakanishi et al., 2017; Cheong et al., 2020):

 $Social sniffing index = \frac{(Social sniffing time)}{(Object sniffing time) + (Social sniffing time)},$

Social interaction index

= (Time in social zone) (Time in object zone) + (Time in social zone)

Resident-intruder test

We devised the resident-intruder (RI) test as previously described (Tan et al., 2021). The mouse was single housed for 24 h and then exposed to an intruder mouse in the home cage for 10 min. We used a slightly smaller (10–20% lighter) unfamiliar sex-matched and strain-matched novel mouse as an intruder. The offensive behaviors of the subject mouse against the intruder, including lateral threat, upright posture, clinch attack, and chase, were scored as aggressive behaviors. In addition, the nonoffensive social behaviors in the RI test, such as nose-to-nose interaction and anogenital sniffing, were scored. All scoring was performed blindly.

Measuring estrous cycle in female mice

We monitored the estrous cycle of female mice using vaginal cytology (McLean et al., 2012). Vaginal smears from subject mice were collected right after the sociability test. Smears were placed on slides and stained with 0.1% crystal violet (catalog #V5265, Sigma-Aldrich). Each estrous cycle stage was determined based on the cell type that was mainly observed in the smear sample among the three types: nucleated epithelial cells, cornified squamous epithelial cells, and leukocytes. To examine the difference in estrous cycles between GH and PWSI mice, we performed the vaginal cytology for 4 consecutive days in each mouse.

In vivo extracellular recordings for measuring sniffing-induced activity

We performed all recordings in the head-fixed and awake mice. We marked the top of OFC_L and right OFC (OFC_R) and implanted customdesigned head plates on the skull of mice using dental cement and small screws. After 2–3 d of recovery, we habituated mice to be head fixed for 20 min/d for 3 d. After the habituation, we performed the recording in the OFC by making a small craniotomy on the marked spots and inserting 32-channel silicon probes (catalog #A1X32-Poly2-10 mm-50–177 or #A1X32-Poly3-10 mm-50-177, NeuroNexus) vertically into each OFC using a microdrive manipulator (Siskiyou). We first recorded spiking activity in the OFC without presenting a stranger mouse for 10 min (object_{sniffing} session) and next continuously recorded neural activity while presenting a head-fixed stranger mouse in front of the recorded mouse for 10 min (social_{sniffing} session). During the recording, we monitored and recorded the sniffing behavior of the mouse by using a web camera (model LifeCam HD-3000, Microsoft). All extracellular signals were recorded at a 30 kHz sampling rate, filtered between 500 and \sim 5000 Hz, and amplified by a miniature digital head stage (CerePlex μ , Blackrock Neurotech). After the recording, we confirmed the recording sites by injecting red retrobeads (Lumafluor) into the recording areas and performing histology. If the recording sites were outside of the OFC, we did not use those recording data in further analysis.

Automatic quantification of sniffing behaviors. We developed a MATLAB code to automatically analyze sniffing events during object-sniffing and social-sniffing sessions in head-fixed mice. First, we converted all the RGB image frames of a recorded video clip to grayscale using the *rgb2gray* function. Next, we chose the ROI, including the nose and whiskers of the mouse being recorded, made a binary matrix with 1 inside the ROI and 0 outside the ROI of each pixel, and multiplied an original image by the binary matrix of each corresponding pixel along all the frames (ROI_1). We subtracted background brightness from ROI_1 to eliminate overall brightness fluctuations (ROI_2) and calculated deviations of brightness within the ROI by subtracting the mean brightness of each pixel (ROI ΔF). To detect sniffing events, we subtracted a moving average (F_{0i} ; sliding window = 50 frames) from ROI ΔF (ROI ΔF - F_0), and set a threshold by multiplying a median of absolute ROI ΔF - F_0 by 4. If ROI ΔF - F_0 in each frame crossed the threshold, we included the frame in a sniffing event.

Neural activity analysis. We performed the spike-sorting using Klusters software (http://neurosuite.sourceforge.net/; Hazan et al., 2006). We grouped 32-channel data into eight groups and sorted single units by principal components analysis. We only isolated single units that show a >2 ms refractory period in their autocorrelogram. We performed all other analyses using custom-made codes in MATLAB (MathWorks). We sorted sniffing trials in object and social sessions by identifying sniffing events across the recording sessions. We only used trials that satisfied the following two criteria: (1) intersniffing interval (time between the offset of sniffing in the current trial and the onset time of the next trial) >5 s; and (2) sniffing duration (time between the onset and the offset of sniffing in one trial) >1 s. To plot the perievent time histogram of all neurons as a population, we performed z-scoring by averaging firing rates (in hertz) of individual units across trials and normalized them by 1 s baseline activity (3 to 2 s before the onset of sniffing). The normalized firing rates were convolved with a Gaussian filter (σ = 50 ms) only for visualization purposes. To identify the neurons showing significant sniffinginduced activity, we statistically compared the firing rates of individual neurons during the 2 s preperiod (3 to 1 s before the onset of sniffing) and 2 s post-period (0 to 2 s after the onset of sniffing) of each sniffing trial by bootstrapping (p < 0.01, n = 5000). We quantified the sniffing-induced activity of neurons by averaging z values from 0 to 2 s after the onset of sniffing. We classified cell types of recorded neurons based on spike waveforms by quantifying spike widths of each isolated neuron (time between peak and through of the spike, in microseconds). If the spike width was <350 or $>450 \ \mu s$, recorded neurons were classified as fastspiking (FS) neurons or regular-spiking (RS) neurons, respectively. For the baseline activity analysis, we quantified the average firing rates (in hertz) from 0 to 2 s before the onset of sniffing.

Perineuronal net degradation

To remove the PNN, we reconstituted protease-free chABC (Amsbio) in PBS to a final concentration of 61 U/ml. We anesthetized mice by inhalation of 1.5% isoflurane in oxygen on the stereotaxic apparatus. We injected 0.5 μ l of PBS or chABC into the left OFC [anteroposterior (AP), +2.35; mediolateral (ML), +1.3; dorsoventral (DV), -2.5] of female mice at P70 to P75. We performed behavior experiments 7 d after the injection and killed the mice to confirm PNN degradation by histology. To stain the PNN in the brain slice containing the OFC_L, we treated the FITC-conjugated WFA (catalog #FL-1351, Vector Laboratories) as a secondary antibody, diluted at 1:500 in 1× PBS, for 2 h at RT before mounting the slices in a transparent mounting medium (catalog #S3023, DAKO). We manually and blindly counted the number of WFA⁺ neurons within 0.5 × 0.5 mm² ROIs in three representative coronal sections covering OFC.

CRISPR/Cas9 experiments

Generating the PX552-sgPV-tdTomato. To generate the AAV-tdTomato (tdTom) vector, we first replaced the eGFP gene in PX552

(catalog #60958, Addgene) with tdTomato. The tdTomato template was obtained from pAAV-FLEX-tdTomato (catalog #28306, Addgene). Next, we cloned two AAV-tdTomato plasmids, both of which induce the targeted mutations at the parvalbumin gene (sgPV). One virus expresses two single-guide RNAs (sgRNAs) targeting parvalbumin exon4: sgRNA1, 5'-AGACAAGTCTCTGGCATCTG-3' and sgRNA2, 5'-AAGGATGGGGA CGGCAAGAT-3' (cloned recombinant DNA fragment: AGACAAGTC TCTGGCATCTGGTTTTAGAGCTAGAAATAGCAAGTTAAAATAAG GCTAGTCCGTTATCAACTTGAAAAAGTGGCACCGAGTCGGTGCA ACAAAGCACCAGTGGTCTAGTGGTAGAATAGTACCCTGCCACG GTACAGACCCGGGTTCGATTCCCGGCTGGTGCAAAGGATGGG GACGGCAAGATGTTTTAGAGCTAGAAATAGCAAGTTAAAATAA GGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCGAGTCGGTGC; Bioneer/in-house PCR amplification), and the other virus expresses two sgRNAs targeting parvalbumin exon5: sgRNA3, 5'-GCTAAGTGGCGCTG ACTGCT-3' and sgRNA4, 5'-GGCCGCGAGAAGGACTGAGA-3'(Cloned recombinant DNA fragment: GCTAAGTGGCGCTGAC TGCTGTTTTAGAGCTAGAAATAGCAAGTTAAAATAAGGCTA GTCCGTTATCAACTTGAAAAAGTGGCACCGAGTCGGTGCAA CAAAGCACCAGTGGTCTAGTGGTAGAATAGTACCCTGCCAC GGTACAGACCCGGGTTCGATTCCCGGCTGGTGCAGGCCGCG AGAAGGACTGAGAGTTTTAGAGCTAGAAATAGCAAGTTAAA ATAAGGCTAGTCCGTTATCAACTTGAAAAAGTGGCACCGAG TCGGTGC; Bioneer/in-house PCR amplification).

After cloning the AAV-tdTomato vectors, we transfected them with packaging plasmid into HEK293T (catalog #HC20005, KCTC) to produce AAV-sgPV-tdTomato (exon4 and exon5) and AAV-tdTomato, using Lipofectamine 2000 (catalog #11668–019, Thermo Fisher Scientific). After 72 h, we collected adeno-associated virus [AAV (AAV2)] particles from the cell lysates and performed iodixanol gradient ultracentrifugation. Next, we concentrated the viral particles with Amicon Ultra Centrifugal Filter (Millipore) and quantified the concentration of viral particles by quantitative PCR (in genome copies/ml: AAV-tdTomato, 3.92×10^{14} ; AAV-sgPV(exon4)-tdTomato, 3.07×10^{12} ; AAV-sgPV (exon5)-tdTomato, 7.17×10^{12}).

Validation of CRISPR/Cas9 in the Neuro2a cell. To identify the efficiency of sgPV in knocking down the PV, we transfected cloned AAV vectors with the Cas9-eGFP-expressing vector (PX458) in the Neuro2a cells. After 72 h, we collected and sorted tdTomato/eGFP double-positive cells using FACSAria II (BD Biosciences). Next, we purified genomic DNA from sorted cells and performed PCR to amplify the target sites of the PV gene, adding the index sequence for sequencing. Last, we subjected the samples to sequencing (MiniSeq System, Illumina) and analyzed insertion and deletion frequencies using CRISPR RGEN Tools (http://www.rgenome.net/).

Behavior and physiological experiments with CRISPR/Cas9 mice. To express Cas9 selectively in PV⁺ neurons, we crossed the PV-Cre mice (catalog #008069, The Jackson Laboratory) with the Rosa26-floxed STOP-Cas9 knock-in mice (catalog #024857, The Jackson Laboratory). We injected tdTom or AAV-sgPV-tdTomato (sgPV) into the crossed mice (PV::Cas9) on P50 to P65. From P78 to P85, we performed the behavior test and *in vivo* recording. To verify whether the AAV expression knocked down the PV effectively, we collected brain samples of the injected mice and performed immunohistochemistry as described above. We used a combination of the primary antibodies (rabbit anti-PV for PV staining, goat anti-GFP for Cas9 staining) and the secondary antibodies (Alexa Fluor 405 donkey anti-goat IgG and Alexa Fluor 647 donkey anti-rabbit IgG were used).

Whole-cell patch-clamp recordings in brain slices. For electrophysiology experiments, tdTom/sgPV-expressed brain samples were collected from anesthetized PV::Cas9 mice with isoflurane (Terrell) and sliced (coronal, $300 \,\mu$ m thickness) using a vibratome (model VT1200, Leica). Brain extraction and sectioning were conducted in ice-cold dissection buffer containing the following (in mM): 212 sucrose, 25 NaHCO₃, 5 KCl, 1.25 NaH₂PO₄, 0.5 CaCl₂, 3.5 MgSO₄, 10 D-glucose, 1.25 L-ascorbic acid, and 2 Na-pyruvate (Pyr), bubbled with 95% O₂/5% CO₂. The sectioned slices were incubated in a recovery chamber at 32°C with normal ACSF (in mM: 125 NaCl, 2.5 KCl, 1.25 NaH₂PO₄, 25 NaHCO₃, 10 glucose, 2.5 CaCl₂, and 1.3 MgCl₂, oxygenated with 95% O₂/5% CO₂). After

60 min of recovery at 32°C, slices were incubated for 30 min at room temperature (20–25°C). The slice was transferred to a recording chamber at 27–29°C with circulating ACSF saturated with 95% O₂ and 5% CO₂. Recording pipettes were made from thin-walled borosilicate capillaries (catalog #30–0065, Harvard Apparatus) with resistance of 3.0–4.0 M\Omega using a two-step vertical puller (model PC-10, Narishige).

Whole-cell patch-clamp recordings were conducted using an amplifier (MultiClamp 700B, Molecular Devices) and digitizer (Digidata 1550, Molecular Devices). We recorded both AAV-expressed and nonexpressed neurons in the OFC_L. We confirmed the AAV expression by imaging the red fluorescence of the patched cells under the fluorescent microscope (Olympus). During the recordings, hyperpolarizing step pulses (5 mV, 40 ms) were injected into the cell to monitor membrane resistance calculated by the peak amplitude of the capacitance current. Data from cells that changed the resistance by >20% were excluded. To measure the intrinsic excitability, recording pipettes $(3.0-4.0M\Omega)$ were filled with an internal solution containing the following (in mM): 137 K-gluconate, 5 KCl, 10 HEPES, 0.2 EGTA, 10 Na-phosphocreatine, 4 Mg-ATP, and 0.5 Na-GTP, at pH 7.2 and 280 mOsm. Picrotoxin (100 μ M; Sigma-Aldrich), NBQX (10 μ M, Tocris Bioscience), and D-AP5 (50 μ M, Tocris Bioscience) were added to ACSF to inhibit postsynaptic responses. After gigaseal and rupture, currents were clamped, and resting membrane potential was maintained at -70 mV. To measure the I-V curve, current inputs were increased from -300 to 120 pA in increments of 30 pA/sweep with a time interval of 2 s. To measure the current-firing (I-F) curve, current inputs were increased from 0 to 660 pA in increments of 60 pA/sweep with a time interval of 10 s. For measuring spontaneous EPSCs (sEPSCs) in the OFC PV^+ neurons, glass pipettes were filled with an internal solution composed of the following (in mM): 100 CsMeSO₄, 10 TEA-Cl, 8 NaCl, 10 HEPES, 5 QX-314-Cl, 2 Mg-ATP, 0.3 Na-GTP, and 10 EGTA, at pH 7.25 and 295 mOsm at a holding potential of -70 mV. Picrotoxin (100 μ M) was added to ACSF to block IPSCs. For measuring spontaneous IPSCs (sIPSCs) in the OFC excitatory neurons, recording pipettes were filled with an internal solution containing the following (in mM): 120 CsCl, 10 TEA-Cl, 8 NaCl, 10 HEPES, 5 QX-314-Cl, 4 Mg-ATP, 0.3 Na-GTP, and 10 EGTA, at pH 7.35 and 280 mOsm holding at -70 mV. NBQX and D-AP5 were added to block EPSCs in the circulating ACSF. Data were acquired by Clampex 10.2 (Molecular Devices) and analyzed by Clampfit 10 (Molecular Devices).

Wireless optogenetic modulation of the OFC neurons in vivo

Design and fabrication of flexible optoelectronic neural probes. Flexible optoelectronic probes were designed using electronic design automation software (Altium Designer 18.0, Altium Limited) and produced using a conventional photolithography process. Neural probes (0.1 mm thick) were designed in a bilateral form, in which the right and the left probes (for each probe: width, 0.38 mm; length, 4 mm) were separated by 2.6 mm to access both regions of the mouse brain. Copper traces and electrodes (18 μ m thick) were patterned on one side (i.e., top layer) of the polyimide substrate (25 μ m thick). In contrast, the other side (i.e., the bottom layer) was covered with a copper layer (18 μ m thick) to stiffen the probe for penetration into brain tissue. The probe tip part, where the microscale inorganic light-emitting diodes (μ -ILEDs; TR2227, Cree; blue, 473 nm; green, 532 nm; $270 \times 220 \times 50 \ \mu m^3$) were attached, was not covered with a copper layer to allow light emission from both sides of the probe. The fabricated probes showed an optical intensity of \sim 405 mW/mm² at the top layer and \sim 45.3 mW/mm² at the bottom layer, which sufficiently exceeds the threshold intensity for channelrhodopsin-2 (ChR2) activation (~1 mW/mm²). A three-pin plug-nplay female pin connector (model M50-3130345, Harwin) and two μ -ILEDs were mounted on the top layer electrodes after applying a lowtemperature solder paste (SMDLTLFP10T5, Chip Quik) and soldered using a precision soldering iron (NASE-2C, JBC Tools) with a temperature of 250°C. After confirming that every component was robustly attached, probes were coated with Parylene C (7 μ m thick) for waterproofing. Last, the probes were vertically bent and attached to a 3Dprinted probe holder, followed by fixing them in place using epoxy (5 Minute Epoxy, Permatex).

Design of programmable wireless control module with detachable configuration. A Bluetooth Low Energy (BLE)-based wireless control module was designed to be easily attached and detached with a bilateral optoelectronic probe and a battery through a plug-n-play connection (Qazi et al., 2019; Lee et al., 2020). This module controls the optical output of the μ -ILEDs through smartphone manipulation. The electronic circuit of the wireless module consists of a rechargeable lithium polymer (LiPo) battery (GM300910-PCB, PowerStream Technology; 12 mAh; $9 \times 10 \times 3$ mm³), a BLE System-on-Chip (SoC; EYSHSNZWZ, Taiyo Yuden; $8.55 \times 3.25 \times 0.3$ mm³), and various other electronic components (a linear voltage regulator, four decoupling capacitors, two resistors, and two indicator LEDs). The current supplied from the LiPo battery is regulated and stabilized by a voltage regulator (NCP4624DMU30TCG, onsemi) and decoupling capacitors and flows to the BLE SoC to operate the module. Two indicator LEDs [model VAOL-S4RP4, VCC; red (624 nm wavelength)] were placed at the output pins of BLE SoC, each connected in series with a resistor (6.04 k Ω ; model RK73H1HTTC6041F, KOA Speer Electronics), to visually show the current operation status of the module during the animal experiment-for example, the indicator LED blinks every 5 s during optical stimulation. The wireless control module can be assembled with optoelectronic probes through three-pin plug-n-play connectors (male: model M50-3630342, Harwin; female: model M50-3130345, Harwin) and assembled with LiPo battery through two-pin connectors (male: model M50-3630242, Harwin; female: model M50-3130245, Harwin). Every component except the LiPo battery was soldered on the wireless module flexible printed circuit board of the wireless module $(9.3 \times 7.6 \times 0.1)$ mm³), which was designed and manufactured through the same method used for the μ -ILED probes. The module was wirelessly programmed over the air (Nordic Device Firmware Update bootloader) to provide a pulse current (20 Hz with 25 ms pulse width; i.e., 50% duty cycle) or continuous current (i.e., 100% duty cycle) to the connected bilateral μ -ILED probes. The BLE-based smartphone app was developed using commercial app development software (Android Studio) and used to control the μ -ILED probes for wireless optogenetics.

Virus injection and probe implantation surgery. For optogenetic activation of excitatory neurons in the left OFC, we injected the AAV2-CAMKIIα-eYFP (enhanced yellow fluorescent protein; catalog #VB1941, Vector Biolabs) or AAV2-CAMKII α -ChR2-eYFP (Addgene) virus in the OFC_{L} (AP, +2.35; ML, ±1.3; DV, -2.5) of female WT mice between P50 and P60. For optogenetic inactivation of PV⁺ neurons in the OFC_L, we injected the AAV9-EF1*a*-DIO-eYFP (UNC Vector Core) or AAV9-EF1α-DIO-eArch3.0-eYFP (UNC Vector Core) virus into the OFC_L of female PV-Cre transgenic mice. For the injection, we anesthetized mice using 1.5% isoflurane in oxygen inhalation and performed stereotaxic surgery. Each injection contained 400 nl of AAV infused at a rate of 0.4 nl/s using a microinjector. We recovered the mice for 7 d after the AAV injection and anesthetized mice again to implant the wireless optogenetic LED probe into the virus-expressing OFC area. Because of the thickness of the probe (0.1 mm) and the location of the LED on the probe (+0.22 mm from the tip), we implanted the probe into the anterior and ventral part of the OFC (AP, ± 2.45 ; ML, ± 1.3 ; DV, -2.77) to target the LED facing the virus-expressing area without any lesion in it.

Behavior test. To connect the small module part to the implanted probe on the mouse head, we briefly anesthetized the mice by isoflurane inhalation. After the module was connected, the mice had 10 min recovery phase in their home cage. On Day 0, we habituated mice wearing a module for 15 min in the open field chamber. On Day 1, we habituated the mice wearing a module in the sociability chamber for 5 min. We then presented a novel social target and a novel object (O) for 5 min and recorded their approaching behaviors to measure the sociability of the mice. On Day 2, we performed a sociability test with blue or green LED stimulation for optogenetic modulation. We gave 20 Hz blue light stimulation by 473 nm LED for activation or continuous green light stimulation by 532 nm LED for inactivation. The LED connected to the probe during the 5 min sociability test session using the custom-designed flexible optoelectronic neural probes. Interaction times of mice were analyzed on Day 1 and Day 2, as described previously. On Day 3, we performed an open field test for 5 min with optogenetic stimulation and 5 min without stimulation to measure the effect of light activation on the locomotion behavior. After the behavior test, we performed the histology of the brain samples from the mice to check the virus expression and probe the implanted site. We collected data only if the virus expressed >65% of the total volume of the OFC and the probe was targeted at the virus-expressed area.

Experimental design and statistical analysis

All statistical analyses were performed using custom codes in IBM SPSS statistics 21 (IBM) and OriginPro 2019 (Origin). The normal distribution of data were verified using the Shapiro–Wilk normality test. For parametric tests, we used the paired Student's *t* test to compare matched pairs, unpaired Student's *t* test to compare two groups, one-way ANOVA with Fisher's least significant difference *post hoc* test to compare multiple groups, and two-way ANOVA to compare *I–F* curves of PV^+ neurons. For nonparametric tests, we used the Wilcoxon signed-rank test to compare matched pairs, the Mann–Whitney *U* test to compare between two independent groups, and the Kolmogorov–Smirnov test to compare the distributions of two groups. The number of animals analyzed by statistical tests and the exact *p* values from each experiment were reported within each figure legend.

Results

Postweaning social isolation enhances sociability in female mice

Previous reports have shown that both short-term and long-term social isolation affects mouse social behaviors (Makinodan et al., 2012; Matthews et al., 2016). However, the effects differ depending on the duration of isolation, and the data were collected only in male mice. To examine whether short-term and long-term social isolation effects on mouse sociability differ between the sexes, we performed the sociability test in both male and female mice. For the sociability test, we put two small chambers at diagonal corners in the maze, each of which contained a novel object (O) or a novel and sex-matched mouse (M), and measured the relative times of mice sniffing or spending at or nearby the novel mouse compared with those times at the novel object as indices of the mouse sociability. We quantified the sociability in mice that had experienced three different postweaning social conditions at P80: GH, SI for 1 d (1dSI), or SI for 60 d after weaning (i.e., PWSI; Fig. 1A,B). Both the sniffing time of 1dSI mice when they came into contact with a novel mouse and their time spent in the social interaction zone did not differ significantly from those of GH mice in both males and females (Fig. 1*C*,*D*).

Notably, PWSI induced changes in sociability in a sex-specific manner: only female mice showed a significant increase in socialsniffing time and social interaction time, while PWSI male mice showed normal sociability, the same as the GH mice (Fig. 1*C*,*D*). We also confirmed that the effect of PWSI on female mice was not simply because of hormonal changes that occur during the estrous cycle. First, we confirmed that the estrous cycle was not different between GH and PWSI female mice (Fig. 1*E*). We next confirmed that the mouse sociability (both social-sniffing index and social interaction index) that we measured was consistent across the estrous cycle in female mice (Fig. 1*F*–*H*). Therefore, our data indicate that long-term social isolation during adolescence affects the social behavior of females more than males, causing hypersociability in females.

PWSI reduces PV expression in the OFC_L of female mice

We next examined whether the social experience during adolescence is required to mature PV^+ inhibitory circuits in the OFC. We quantified the number of PV^+ or PNN^+ neurons in the OFC of the GH and the PWSI mice at P80. We collected data from male and



Figure 1. PWSI during adolescence enhances social approaching behavior in female mice. *A*, Experimental design of the sociability test. Schematic of GH, 1dSI, and PWSI. Note that the sociability behavior was measured at P80. *B*, Representative heatmaps indicating mouse locations in the square chamber during the sociability test. In the chamber, a container containing a novel object (0) and a container containing a novel mouse (M) were located at diagonal corners. Heatmaps were calculated in EthoVision XT as the mean duration of time spent in an area for each group. *C*, The social-sniffing index displays the social sniffing times divided by the total sniffing times. One-way ANOVA: female: $F_{(2,101)} = 8.431$, ***p < 0.001; male, $F_{(2,36)} = 0.015$, p = 0.985, Fisher's least significant difference *post hoc* test (female: GH vs PWSI, ***p < 0.001; 1dSI vs PWSI, **p = 0.009; GH vs 1dSI, p = 0.967). Note that only the PWSI female mice showed a significant increase in social-sniffing behavior. *D*, The social interaction index displays the time spent in the social zone divided by the total time spent in both object and social interaction zones. One-way ANOVA: female: $F_{(2,101)} = 3.496$, *p = 0.034; male: $F_{(2,36)} = 0.109$, p = 0.897, Fisher's least significant difference *post hoc* test (female, GH vs PWSI, *p = 0.027; 1dSI vs PWSI, *p = 0.031; GH vs 1dSI, p = 0.494). Note that only the PWSI female mice showed a significant increase in social interaction behavior. *E*, The number of estrous cycles per 4 d of GH (n = 5) and PWSI mice (n = 5). The duration of estrous cycles was not different between groups. Mann–Whitney U test, p = 0.912. *F*, Representative images of cells in the vaginal smear that were stained with crystal violet. The estrous cycles of female mice series across the estrous cycle. One-way ANOVA: $F_{(3,30)} = 0.513$, p = 0.677. All data are presented as the mean \pm SEM. For *C* and *D*: Female: GH, n = 54 mice; 1dSI, n = 14 mice; PWSI, n = 36 mice;

female mice separately. We first immunostained both for the PV, a marker for fast-spiking GABAergic neurons, and the NeuN, a marker for all neurons (Fig. 2*A*). We counted the number of immunostained neurons automatically in the three coronal sections that were collected across the center of the OFC (+2.1 to approximately +2.34 mm from the bregma) per each hemisphere in each mouse (see Materials and Methods; Fig. 2*B*,*C*). Interestingly, the number of PV⁺ and NeuN⁺ neurons in the OFC was significantly lower in the brains of PWSI female mice compared with the GH female mice, especially in the left hemisphere (Fig. 2*D*–*F*). In male

mice, however, the number of PV^+ and $NeuN^+$ neurons was not significantly different between GH and PWSI groups (Fig. 2D-F). When we counted the number of PV^- and $NeuN^+$ neurons by subtracting the number of PV^+ neurons from the total number of NeuN⁺ neurons, there was no difference between groups (Fig. 2F, filled bars; female, OFC_L , p = 1, Mann–Whitney U test). These data indicate that the reduction of NeuN⁺ neurons preferentially occurs in the PV^+ neurons rather than in overall PV^- neurons in the OFC_L (Fig. 2F). We next quantified the PNN⁺ neurons in the OFC by staining for WFA, a marker for PNNs



Figure 2. PV expression in the OFC₁ decreases in the PWSI female mice. A, A representative image of the immunostained PV and NeuN in the OFC of a mouse brain coronal section marked with the ROIs (0.09 mm²). Scale bar, 1 mm. B, Validation of the automatic cell-counting method. Left, Representative images of the NeuN (white) and PV (red) immunostained in the OFC1 of GH female mice. The left panels show original images used for manual counting, and the right panels show automatically detected cell boundaries from original images. The bottom numbers indicate the counted cell numbers. Right, The average numbers of NeuN⁺ and PV⁺ neurons in the OFC_L of GH female mice from the manual counting (gray) and automatic counting (black; n = 6). Note that there is no difference between numbers from manual and automatic counting. Mann–Whitney U test: NeuN, p = 0.200; PV, p = 1.000. C, Schematic brain images showing the selection of brain slices for cell counting. Three brain slices were selected between AP +2.1 and +2.34 mm, with an interval of 60 μ m. **D**, Representative fluorescence images of the PV (red) and the NeuN (white) immunostained in the OFC₁ and OFC_{R} of P80 GH and PWSI female mice. Scale bars, 100 μ m. *E*, The number of PV⁺ neurons in the OFC. PWSI decreased the number of PV⁺ neurons in the female OFC. Mann–Whitney U test: female left, *p = 0.030; female right, p = 0.297; male left, p = 0.623; male right, p = 1.000. F, The number of NeuN⁺ neurons in the OFC. Mann– Whitney U test: female left, *p = 0.024; female right, p = 1.000; male left, p = 0.903; male right, p = 1.000. Slashed bars indicate the number of PV⁺ (NeuN⁺) neurons and filled bars indicate PV^- (NeuN⁺) neurons. Note that PWSI female mice showed a selective reduction in PV^+ (NeuN⁺) neurons in OFC₁. G, A representative image of a mouse coronal brain section that was immunostained for PV (red) and WFA (green). Scale bar, 1000 μ m. H, Representative fluorescence images of the OFC₁ immunostained with anti-PV and anti-WFA antibodies. I, The number of WFA $^+$ neurons in the OFC left and right of male and female mice. Mann–Whitney U test: GH vs PWSI: female left, p = 0.46; female right, p = 0.185; male left, p = 0.337; male right, p = 0.42. All data are presented as the mean \pm SEM. PWSI, Mice isolated from P21 to P80. For **E** and **F**: female: GH, n = 6 mice; PWSI, n = 6 mice; male: GH, n = 4 mice; PWSI, n = 5 mice. For H-J: female: GH, n = 12 mice; PWSI, n = 11 mice; male: GH, n = 7 mice; PWSI, n = 7 mice. *p < 0.05. n.s., Not significant.

(Fig. 2*G*). Both PV^+ and PV^- neurons were surrounded by the PNNs in the OFC (Fig. 2*H*). We found that the numbers of WFA⁺ neurons in the OFC were not different between GH and PWSI in both male and female mice (Fig. 2*I*). Collectively, our data indicate that social experience during adolescence is crucial for the maturation of PV^+ neurons, especially the PV expression, in the OFC_L of female mice.

Late adolescence is a critical window for maturing sociability in female mice

PWSI mice experienced long-term social isolation from when they were weaned at P21 to when they became adults at P80. This period includes both early juvenile periods before the onset of puberty and late adolescence after the onset of puberty. Since puberty leads to sexually dimorphic changes in behaviors, we



Figure 3. Social isolation during late adolescence causes hypersociability only in female mice. A, The social-sniffing indices of GH female and male mice. The sociability test was performed at P50 and P80 in the same mouse. Paired: Student's t test within a group: female, **p = 0.008; male, p = 0.756. Unpaired: Student's t test between groups: P50, p = 0.285; P80, p = 0.731. Female, n =11 mice; male, n = 8 mice. B, Left, Representative fluorescence images of the OFC₁ immunostained with anti-PV antibodies (red). Brain sections were collected from P21 (top), P50 (middle), and P80 (bottom) GH female mice. Right, The number of PV⁺ neurons in the OFC_R of female mice. One-way ANOVA: female OFC₁: F_(2,22) = 8.473, **p = 0.001, Fisher's least significant difference post hoc test (P21 vs P50, p = 0.580; P50 vs P80, **p = 0.003; P21 vs P80, ***p < 0.001); female OFC_R: $F_{(2.28)} = 2.856$, *p = 0.010, Fisher's least significant difference post hoc test (P21 vs P50, p = 0.204; P50 vs P80, p = 0.065; P21 vs P80, **p = 0.003). Female: P21, n = 9 mice; P50, n = 11 mice; P80, n = 15 mice. C, Schematic of GH, P50SI, and re-GH at P50 after social isolation at P21. Note that the behavioral tests were performed at P80. **D**, The social-sniffing indices of GH, P50SI, and re-GH mice. One-way ANOVA, female: F_(2,44) = 5.069, *p = 0.01, Fisher's least significant difference post hoc test (female: GH vs P50SI, **p = 0.007; GH vs re-GH, p = 0.276; P50SI vs re-GH, p = 0.127). Mann–Whitney U test, male: GH vs P50SI, p = 0.505. Note that only the P50SI female mice showed a significant increase in social-sniffing behavior compared with GH female mice. Female: GH, n = 23 mice; P50SI, n = 13 mice; re-GH, n = 11 mice; male: GH, n = 10 mice; P50SI, n = 8 mice. *E*, The number of PV⁺ neurons in the OFC of GH (black) and re-GH (blue) female mice. Unpaired, Student's t test: OFC left, p = 0.873; OFC right, p = 0.120. Female: GH, n = 15 mice; re-GH, n = 7 mice. F, The RI test (RIT) was performed for P50SI and GH female and male mice at P80. Top, A representative image of RIT. Bottom, The list of analyzed behaviors. The offensive behavior included lateral threat, upright posture, clinch attack, and chase. The social behavior included nose to nose (N-N) interaction and anogenital sniffing. G, The RIT results of GH and P50SI female mice. Left, Duration of offensive behaviors. Mann–Whitney U test, p = 0.699. Right, Duration of social behaviors. Mann–Whitney U test: N-N interaction, ***p < 0.001; anogenital sniffing, **p = 0.008. Female P50SI mice showed a significant increase in social behaviors without any change in offensive behaviors. H. The RIT results of GH and P50SI male mice. Left, Duration of offensive behaviors. Mann–Whitney U test, *p = 0.037. Right, Duration of social behaviors. Mann–Whitney U test: N-N interaction, p = 0.095; anogenital sniffing, p = 0.835. Male P50SI mice showed a significant increase in offensive behaviors without any change in social behaviors. All data are presented as the mean \pm SEM. For *F*–*H*: female: GH, *n* = 11 mice; P50SI, *n* = 11 mice; male: GH, *n* = 5 mice; P50SI, *n* = 5 mice. ****p* < 0.001, ***p* < 0.05; n.s., Not significant.

wondered how mouse sociability matures after the onset of puberty in males and females. We thus tracked sociability changes from P50 (late adolescence; Laviola et al., 2003) to P80 (adult) in both sexes of GH mice by measuring the social-sniffing index at P50 and at P80. Interestingly, we found a significant decrease in sociability at P80 in female mice but not in male mice (Fig. 3*A*). We further found that during this period, PV^+ neurons increased significantly in female OFC_L. In contrast, OFC_R did not show such



Figure 4. PWSI reduces sniffing-induced activity in the OFC_L of female mice. *A*, Left, Experimental design for *in vivo* multichannel recording of the neurons responsive to social sniffing in the OFC of head-fixed mice. Right, A representative top view image for video analysis. The blue circle indicates the ROI used to analyze the brightness changes in the video clip. *B*, Schematic illustration of mouse brain sections marked with the recording sites in the OFC of GH or PWSI female mice. Blue shading, OFC; colored bars, recording sites confirmed by *post hoc* histology. *C*, Representative sniffing events measured in GH female mice. Gray trace, Raw brightness changes in the ROI (ROI ΔF); green trace, moving average in the ROI (F_0 ; sliding window, 50 frames); blue and red trace, normalized brightness changes in the ROI ($\Delta F - F_0$); circles, sniffing events were automatically detected by a MATLAB-based image analysis code; horizontal lines, sniffing events manually detected by an observer in the EthoVision XT software. Note that automatic counting methods detect sniffing events at similar times identified by manual counting. Blue,

an increase between P50 and P80 (Fig. 3*B*). These data suggest that late adolescence from P50 to P80 is a critical window for the maturation of sociability with an increase in PV^+ neurons in the OFC_L in female mice.

We next examined the effect of social isolation during late adolescence in both female and male mice. We directly isolated the mice after P50 until P80 (P50SI; Fig. 3C). In addition, we regroup housed (re-GH) a group of PWSI female mice at P50 to make them re-experience social interaction during late adolescence (re-GH; Fig. 3C). The P50SI female mice showed an increase in sociability like the PWSI female mice, while neither re-GH female mice nor P50SI male mice showed any changes in sociability compared with control GH mice (Fig. 3D). Not only our modified three chamber-like test, but a standard three-chamber test also showed an increase in sociability in P50SI female mice (data not shown). In addition, the number of PV^+ neurons in re-GH female mice was not significantly different from that in GH female mice in both hemispheres of the OFC (Fig. 3E), indicating that social experience during late adolescence was sufficient to mature the PV^+ neurons in the OFC_L.

Finally, we wondered whether the enhanced sociability in P50SI female mice, measured by our sociability test, was related with changes in aggression. To examine this, we performed the resident-intruder test (Tan et al., 2021) on P50SI male and female mice. We counted both offensive behaviors, such as threatening and attacking behaviors, and social behaviors, such as nose sniffing and anogenital sniffing, separately during the test (Fig. 3*F*). In female mice, the aggression level did not increase by P50SI. Still, the social-approaching behaviors significantly increased (Fig. 3*G*). On the other hand, P50SI male mice showed a significant increase in aggression without any changes in social behaviors (Fig. 3*H*). Collectively, our data indicate that social isolation during late adolescence induced hypersociability

in female mice without increasing aggression, whereas it induced hyperaggression in male mice.

PWSI disrupts social sniffing-induced activity in the OFC_L of female mice

Because fewer PV^+ neurons in the OFC_L were found following PWSI in female mice, we investigated whether the OFC_L neural response to social contact has been altered in head-fixed female mice by in vivo multichannel recordings (Fig. 4A,B). During the recording, both GH and PWSI female mice showed sniffing behaviors toward an object or another mouse under the headfixed condition, and the number and duration of sniffing times did not differ between groups (Fig. 4C,D). We identified neurons that showed significant sniffing-induced activity in the OFC (Fig. 4E-H). Interestingly, in GH females, more neurons showed sniffing-induced activity in the OFC_L than in the OFC_R responses, and their responses were stronger when a mouse sniffed another mouse than when it sniffed an object (Fig. 4E,F). More importantly, these socially responsive neurons in the OFC_L of female mice largely disappeared after PWSI (Fig. 4E,G,H). On the other hand, the sniffing-induced activity of the OFC_R did not differ between GH and PWSI female mice (Fig. 4F). Among the responsive neurons in the OFC_L, the fraction of the fast-spiking neurons was smaller in the PWSI mice than in the GH mice. These data indicate that PWSI decreased social sniffing-induced activity in the OFC_L neurons, particularly including fast-spiking neurons, and reduced overall social representation in the OFC_L.

PV^+ expression, not PNNs in the OFC_L, shapes social behavior in female mice

As we found that PWSI impaired the PV expression in the OFC_L and increased social behavior, we next examined the causal relationship between the reduction of PV expression in the OFC_L and the increase in the sociability of female mice. To reduce the PV expression in vivo, we used the CRISPR/Cas9 system (Swiech et al., 2015) and designed two pairs of sgRNAs to target the fourth and fifth exons of the PV gene (sgPV). We first confirmed that those sgRNAs knocked out the PV gene by either deletion or insertion of the targeted sequences in >85% of the Cas9-transfected Neuro2a cells (Fig. 5A). We then constructed the AAV vectors to deliver those sgRNAs together with the tdTomato gene to neurons in vivo (Fig. 5A) and injected the AAVs into the cortex of PV::Cas9-eGFP mice, which were created by crossing PV-Cre with Cas9^{fl/fl}-eGFP mice (sgPV; Fig. 5B,C). As a control, we injected the AAV that expressed only the tdTomato without sgRNAs into the crossed mice (tdTom). In the sgPV mice, originally PV⁺ neurons were expressed with both Cas9-eGFP and tdTomato, and their PV gene was knocked out by Cas9 in the PV^+ neurons selectively (Fig. 5B,C).

To knock down PV in the OFC_L during late adolescence until mice became adults, we injected the AAVs into the OFC_L when mice were at approximately P50 and examined changes in the sociability of the injected mice at P80. We confirmed the reduction of PV expression in the eGFP⁺/tdTomato⁺ neurons without any decrease in PNN expression by immunostaining the PV and the PNNs in the OFC_L (Fig. 5*D*,*E*). The female mice with PV knockdown in the OFC_L showed a significant increase in sociability (Fig. 5*H*), similar to the PWSI or P50SI females (Figs. 1*B*–*D*, 3*D*). When we knocked down the PV in the OFC_R of female mice or the OFC_L of male mice, we did not see such an increase in sociability compared with the control mice (Fig. 5*H*). We further examined whether a reduction in PNNs surrounding PV⁺ neurons affects female sociability. We infused chABC, an enzyme that

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Object sniffing; red, social sniffing. **D**, Left, Sniffing events automatically counted. Dots, Individual data; lines, mean \pm SEM. Wilcoxon signed-rank test within a group: object versus social: GH, p = 0.465; PWSI, p = 1.000. Mann–Whitney U test between groups: GH versus PWSI: object, p = 0.857; social, p = 0.629. Right, The sniffing events manually identified. Dots, Individual data; lines, mean \pm SEM. Wilcoxon signed-rank test within a group: object versus social: GH, p = 0.686; PWSI, p = 0.109. Mann–Whitney U test between groups: GH versus PWSI: Object, p = 0.393; Social, p = 1.000. GH, n = 4 mice; PWSI, n = 3 mice. **E**, Neural responses associated with sniffing in the female OFC₁. The large pie charts, the OFC₁ neurons showing social-sniffing responses (Social, red), object-sniffing responses (Object, blue), or both social and object sniffing responses (Both, green). Inc and dec, neurons showing significantly increased and decreased activities after the onset of sniffing, respectively (p < 0.01, bootstrap); No resp, Neurons that show no responses. The small pie charts, the waveforms of the responsive (right top) and nonresponsive (left bottom) OFC, neurons classified according to the waveform width. US, Unclassified neurons with a waveform width between 350 and 450 ms. Heatmaps, z-scored activity of all neurons normalized by the presniffing activity. GH, n = 5 mice, n = 84 units; PWSI, n = 3 mice, n = 57 units. **F**, The same as **C**, but for the OFC_R neurons. GH, n = 4 mice, n = 74 units; PWSI, n = 4 mice, n = 60 units. G, Representative raster plots and perievent time histograms of example "Social Inc" OFC neurons from the GH female mouse (top, GH A #5) and from the PWSI female mouse (bottom, SI B #9) aligned by the onset of object sniffing (left, blue) and the onset of social sniffing (right, red). H, Averaged sniffing-induced activity for 2 s (z score) of the top 20% of OFCL neurons that show either increased or decreased responses after the onset of social sniffing. Bars, Mean \pm SEM. Wilcoxon signed-rank test for each bar: Object_{sniffing}, Social_{sniffing}: ***p < 0.001, ***p < 0.001 for GH Increased; p = 0.083, **p = 0.004 for PWSI Increased; p = 0.478, ***p < 0.001 for GH Decreased; *p = 0.045, **p = 0.004 for PWSI Decreased. Wilcoxon signed-rank test within a group: $Object_{sniffing}$ versus $Social_{sniffing}$: *p = 0.047 for GH Increased; *p = 0.023 for PWSI Increased; **p = 0.002 for GH Decreased; p = 0.120 for PWSI Decreased; Mann–Whitney *U* test between groups: Object_{sniffing}, Social_{sniffing}: *p = 0.019, ***p < 0.001 for Increased activity; p = 0.452, p = 0.110 for Decreased activity. GH, n = 17units; PWSI, *n* = 11 units. ****p* < 0.001, ***p* < 0.01, **p* < 0.05; n.s., Not significant.



Figure 5. CRISPR/Cas9-mediated knockdown of PV, but not PNN degradation, in the OFC_L enhances sociability in female mice. *A*, Top, Schematic illustration of the AAV-sgPV-tdTomato expression vector. The sgRNA vector contains the encoding sequence of tdTomato protein for identifying transduced neurons. We used two types of AAVs to target both exon 4 and exon 5 of the PV gene. Bottom, DNA insertion (blue) and deletion (red) mutation ratio (Indel frequency) by the designed sgPVs targeting exon 4 and exon 5 of the PV gene and cloned in the AAV-tdTomato or AAV-sgPV-tdTomato. Scale bars, 20 μ m. Red, tdTomato expressed by AAV vectors; green, Cas9-eGFP transgenically expressed in PV⁺ neurons; blue, immunostained PV proteins. Solid arrows Triple-positive cells; dashed arrows, double-positive and PV⁻ cells. *C*, Quantification of PV⁺ cells among eGFP and tdTomato double-positive cells in the retrosplenial cortex (RSC), injected with AAVs that deliver tdTom or sgPVs, compared with the Cas9-eGFP expression in PV:: Cas9 mice. Numbers in columns indicate the number of brain samples analyzed. One-way ANOVA: $F_{(3,63)} = 6.478$, ***p < 0.001; Fisher's least significant difference *post hoc* test. *D*, CRISPR/Cas9-mediated knockdown of PV in the female OFC_L. Left and middle, Representative fluorescence images of the OFC_L of the female PV::Cas9-eGFP mice injected with the AAV-tdTomato (top, tdTom) or AAV-sgPV-tdTomato (bottom, sgPV). Red, tdTomato; green, Cas9-eGFP; blue, immunostained PV. Scale bars, 50 μ m. Right, A representative image of the OFC_L of a PV::Cas9-eGFP

degrades PNNs by removing sulfate side chains of proteoglycans, into the OFC_L (Fig. 5*F*). Treatment with chABC significantly decreased the number of WFA⁺ neurons in the OFC_L without affecting PV expression (Fig. 5*G*). Not like the female mice knocked down with PV, the female mice treated with chABC showed normal sociability (Fig. 5*I*). Together, our data indicate that the PV expression, but not the PNN maturation, in the OFC_L was a key factor shaping the social behavior of female mice.

Decreased PV expression reduces social sniffing-induced activity via hyperexcitation of OFC_L neurons

We next investigated whether the decrease of PV expression in the OFC_L was responsible for the reduced sniffing-induced activity in the OFC_L of the PWSI female mice. We performed in vivo recordings in female mice injected with sgPV AAVs (sgPV) or injected with tdTomato AAVs (tdTom) in their OFC_L (Fig. 6A-G). First, we found that the average baseline firing rates were significantly higher in the spPV group compared with the tdTom group (Fig. 6D,E). Furthermore, similar to PWSI, knocking down PV by sgPV decreased social sniffing-induced activity in the OFC_L of female mice (sgPV; Fig. 6F,G). Conversely, the expression of tdTomato in the OFC_L did not alter the sniffing-induced activity of female mice (tdTom; Fig. 6F,G). Knocking down PV in the OFC_L did not affect the sniffing behavior itself, as the number and duration of sniffs in the altered mice were similar to those in the tdTom control mice (Fig. 6C). These data indicate that the knockdown of PV increased overall firing activity while reducing the sniffinginduced activity in the OFC_L.

As we observed an increase in the baseline activity of OFC_L neurons by knocking down the PV, we next examined whether the knockdown of PV has altered the excitation–inhibition balance in the OFC_L of female mice. We first performed patch-clamp recordings of the PV⁺ neurons in the OFC_L by targeting tdTomato⁺ neurons of PV::Cas9 mice injected with sgPV or tdTom AAVs into the OFC_L (Fig. 6*H*–*L*). We found that the knockout of the PV gene by sgPV in PV⁺ neurons reduced the intrinsic excitability of those neurons (Fig. 6*I*) without any changes in sEPSCs (Fig. 6*L*). Other membrane properties, such as the half-width of spike waveforms and the input resistance, were normal in those neurons (Fig. 6*J*,*K*). We next measured the sIPSCs in the OFC_L neurons, which were not labeled with the tdTomato in the sgPV and the tdTom mice. These neurons were regular spiking and thus putative excitatory neurons in the OFC_L (Fig. 6*H*,*M*). The OFC_L excitatory neurons in the sgPV mice showed a significant decrease in both amplitudes and frequencies of spontaneous IPSCs compared with the tdTom control mice (Fig. 6*M*). These data indicate that the decreased expression of PV in the OFC_L reduced the excitability of the PV⁺ neurons and, in turn, reduced inhibition of the neighboring neurons in the OFC_L of female mice.

Our recording data suggested that the decrease in inhibition on knocking down the PV gene caused an increase in excitation, which led to a rise in the baseline activity of OFC_L and disruption in sniffing-induced activities (Fig. 6D-G). We thus examined whether the abnormal increase of OFC_L activity caused hypersociability in female mice. We expressed the ChR2 in the excitatory neurons of the OFC_L by injecting the AAV virus, which expresses the ChR2 fused to the eYFP under the CaMKII α promotor, to the wild-type female mice (Fig. 7A,C). We then delivered the light stimulation to the OFC_L by inserting a wireless optogenetic device that we developed (Kim et al., 2021; Qazi et al., 2022) to provide the 20 Hz blue lights during the sociability tests (Fig. 7B; see Materials and Methods). The stimulation was given on the second day of the sociability tests. We measured the first-day sociability (Day 1) without the light stimulation and did not observe any differences in the sociability between the ChR2-eYFP and the eYFP groups. We tested their sociability again on the second day with the blue-light stimulation on the OFC_L (Day 2). The eYFP control mice showed a significant decrease in their sociability on Day 2 compared with Day 1, potentially because of habituation (Fig. 7D,E). On the other hand, the ChR2-eYFP-expressed mice showed higher levels of social-sniffing behaviors with light stimulation than the control mice on Day 2 (Fig. 7D). These data indicate that the optogenetic activation of the OFC_L induced higher sociability and almost no habituation in social approaching behaviors on Day 2 (Fig. 7D,E). We next optogenetically inactivated the PV⁺ neurons in OFC_L directly during the sociability test on Day 2 by expressing the archaerhodopsin-3 (eArch3.0) in PV⁺ neurons and delivering 532 nm light through our wireless probe (Fig. 7G). Similar to the optogenetic activation of excitatory neurons in the OFC_L, the inhibition of PV⁺ neurons induced hypersociability in female mice without habituation on Day 2, when we gave the light stimulation in the OFC_L (Fig. 7H,I). The light stimulation did not affect the locomotion in both groups (Fig. 7F,J). Collectively, our data demonstrate that hyperexcitation of the OFC_L by reducing inhibition caused hypersociability in female mice.

Discussion

Our study found that adolescent social experience is critical for shaping sociability in female mice. Social isolation during late adolescence enhanced sociability in female mice while impairing the maturation of PV^+ neurons and decreasing social sniffinginduced activity in the OFC_L. The knockdown of PV in the OFC_L of female mice induced the same phenotype as the PWSI mice: enhanced sociability and decreased sniffing-induced activity in the OFC_L. These effects were because of the imbalance between excitation and inhibition, as the PV knockdown caused hyperexcitation in the OFC_L. We further found that optogenetic activation of excitatory neurons or suppression of PV⁺ inhibitory neurons in the OFC_L led to enhanced sociability in female mice. This finding supports the idea that the excitation–inhibition balance in the prefrontal cortex is critical for shaping social

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mouse injected with the AAV-sgPV-tdTomato. Scale bar, 1000 μ m. **E**, The number of PV⁺ and WFA⁺ neurons in the OFC₁ injected with the AAV-tdTomato (gray bars) or with the AAV-sqPV-tdTomato (red bars). Mann–Whitney U test: PV, *p = 0.022; WFA, p = 0.306. Female: n = 5 mice for tdTom; n = 6 mice for sgPV. **F**, chABC-induced PNN degradation in the OFC1 of female mice. Left and middle, Immunostained PV (red) and WFA (green) in the OFC₁ of female mice injected with PBS (top) or chABC (bottom). Scale bars, 50 μ m. Right, Representative fluorescence images of the PNN stained with the WFA. Note that the chABC injection selectively degraded PNNs in the OFC_L. Scale bars, 1000 μ m. G, The number of PV^+ and WFA⁺ neurons in the OFC₁ injected with the PBS (gray bars) or the chABC (green bars). Mann–Whitney U test: PV, p = 0.609; WFA, *p = 0.017. Female: n = 7 mice for PBS; n = 7 mice for chABC. *H*, Social-sniffing indices of PV::Cas9 female mice injected with tdTom or sqPV. Mann–Whitney U test: tdTom versus sqPV: female OFC₁, *p = 0.021; female OFC_R, p = 0.885; male OFC₁, p = 0.914. tdTom: female OFC₁, n = 8 mice; female OFC_R, n = 4 mice; male OFC_L, n = 6 mice; sgPV, red: female OFC_L, n = 10 mice; female OFC_R, n = 4 mice; male OFC_{L} , n = 4 mice. I, Social-sniffing indices of mice injected with PBS (gray) or chABC (green). Mann–Whitney U test, p = 0.878. PBS, n = 8 mice; chABC, n = 8. All data are presented as the mean \pm SEM. ***p < 0.001, **p < 0.01, *p < 0.05. n.s., Not significant.



Figure 6. Knockdown of PV reduces sniffing-induced activity and decreases synaptic inhibition in the female OFC₁. A, Schematic illustration of *in vivo* multichannel recording in a head-fixed mouse. B, Schematic illustration of coronal brain sections of in vivo recording sites in the OFCL where tdTomato (tdTom)-expressing or PV sgRNA (sgPV)-expressing AAVs were injected. C, Sniffing events automatically counted for tdTom and sgPV mice. Left, Wilcoxon signed-rank test for within group: tdTom, p = 0.463; sgPV, p = 0.892. Mann–Whitney U test between groups: Object, p = 0.537; Social, p = 0.126. **D**, The cumulative distribution function of baseline firing rates in tdTom and sgPV mice. Baseline firing rates were calculated as the mean firing rates for 2 s before the sniff onset. Kolmogorov–Smirnov test, **p = 0.006. *E*, Mean baseline firing rates of tdTom and sgPV mice. Mann–Whitney U test, ***p < 0.001. *F*, Neural responses associated with the sniffing in the OFC₁ of tdTom-injected (top) or sqPV-injected (bottom) PV::Cas9 female mice. Pie charts, the OFC₁ neurons showing social-sniffing responses (Social, red), object-sniffing responses (Object, blue), or both social-sniffing and object-sniffing responses (Both, green). Inc and dec, neurons showing significantly increased and decreased activities after the onset of sniffing, respectively (p < 0.01, bootstrap); No resp, neurons that show no responses. Heatmaps, z-scored activity of all recorded neurons by the presniffing activity. Note that the number of sniffing-responsive neurons and social sniffing-induced activity was reduced in sgPV females. Heat maps, z-scored activity of all neurons normalized by the presniffing activity. G, Averaged sniffing-induced activity for 2 s (z score) of the top 20% of OFC_L neurons that show either increased or decreased responses after the onset of social sniffing. Wilcoxon signed-rank test for each bar: Object_{sniffing}: Social_{sniffing}: ***p < 0.001, ***p < 0.001 for tdTom Increased; p = 0.154, **p = 0.006 for sgPV Increased; **p = 0.007, ***p < 0.001 for tdTom Decreased; p = 0.053, ** p = 0.006 for sgPV Decreased. Note that the increased number neurons showed larger activity changes after social sniffing than after object sniffing. Wilcoxon signed-rank test within a group: Object_{sniffing} versus Social_{sniffing}: ***p < 0.001 for tdTom Increased; **p = 0.008 for sgPV Increased; p = 0.141 for tdTom Decreased; *p = 0.011 for sgPV Decreased. The social sniffinginduced activity was larger in tdTom females than in sgPV females. Mann–Whitney U test between groups: Object_{sniffing}, Social_{sniffing}, p = 0.067, **p = 0.004 for Increased activity; p = 0.720, p = 0.073 for Decreased activity. tdTom, n = 24 units, sgPV, n = 12 units. H, Physiologic properties of PV⁺ neurons expressed with tdTom or sgPV. Schematic illustration of the whole-cell patch-clamp recordings of PV⁺ neurons (red) or excitatory neurons (blue) in the PV::Cas9 mice injected with the AAVs expressing either tdTom or sgPV-tdTom into the OFCL. *I*, SgPV expression

behaviors in mice (Yizhar et al., 2011). Our data also suggest that adolescent social isolation influences male and female mice differently, both in how the prefrontal cortex matures and social behavior matures.

Importance of social experience during adolescence in shaping female social behavior

Social experience during adolescence is important for the maturation of the mammalian brain (Spear, 2000; Fuhrmann et al., 2015). In humans, social isolation during adolescence can increase the occurrence of neurologic disorders and cause defects in brain function (Almeida et al., 2021). Similarly, depriving rodents of social experience during adolescence caused defects in the maturation of their brains and alterations in social and cognitive behaviors (Cooke et al., 2000; Schubert et al., 2009; Makinodan et al., 2012; Hinton et al., 2019; Potrebic et al., 2022). Furthermore, behavioral and physiological changes caused by adolescent social isolation were often sex specific (Pisu et al., 2016; Bicks et al., 2020; Tan et al., 2021). In this study, we found that PWSI, particularly after puberty (from P50 to P80), impairs the maturation of inhibitory circuits in the OFC_L and causes maladaptation in social exploration behavior in female mice but not in male mice. Our study is unique compared with recent studies that examined the effects of social isolation during early adolescence (Bicks et al., 2020; Tan et al., 2021). Our findings emphasize the importance of later adolescence when females adjust and shape their social-approaching behaviors with the maturation of PV^+ neurons in the OFC_L.

A recent study has shown that the 1dSI increased sociability in male mice (Matthews et al., 2016). We also found that male mice that had experienced 1dSI spent more time interacting with a novel mouse than did GH male mice (GH, 221.11 \pm 17.93 s; 1dSI, 317.97 \pm 31.46 s; p = 0.023, Student's t test). However, the 1dSI male mice also showed a tendency of increased interaction with an object in the other container (GH, 107.29 \pm 9.88 s; 1dSI, 138.28 \pm 23.36 s; p = 0.256, Student's t test). As a result, their social-sniffing and social interaction indices, which indicate the ratio between social interaction and object interaction times, did not change in 1dSI male mice (Fig. 1*C*,*D*). Our data suggest that 1dSI may enhance object exploration behaviors in male mice. This possibility was not tested in the previous report, as they used an empty chamber without any object as opposed to the social chamber. Future studies are required to understand the effect of 1dSI in male mice fully. Regardless, our study focuses on the social isolation effect on female mice, and we found no effects of 1dSI on social and object interaction times in female mice. Our data demonstrate that long-term social isolation during late adolescence affects social behaviors more effectively than the 1dSI in females.

PWSI not only deprives sensory experience from social interaction but also enhances stress in social mammals like mice (Fone and Porkess, 2008). It has been shown that both sensory deprivation and stress can impair the maturation of PV⁺ neurons in the sensory cortex as well as in the hippocampus (Huang et al., 1999; Czeh et al., 2005; Hu et al., 2010). The OFC, which is classified as a secondary olfactory cortex, is one of the key areas involved in olfactory processing (Lipton et al., 1999) and sniffing behaviors (Kareken et al., 2004). Furthermore, the olfactory system is the most important way through which mice detect other mice (Radyushkin et al., 2009). The OFC has a strong connection with the olfactory system and the hippocampus (Wikenheiser and Schoenbaum, 2016). An fMRI study in humans showed that in response to odorant cues with emotional valence, only females showed increased activity in their OFC_L (Royet et al., 2003). Future studies are required to unravel which types of social olfactory cues and stress-related factors affect the maturation of PV⁺ neurons in the OFC_L and social behavior.

Our data suggest that socially evoked activity in the OFC_L is important for reducing social-approaching behaviors when the animal becomes an adult. Excessive sociability is a symptom of patients with Williams–Beuren syndrome (Mimura et al., 2010; Sakurai et al., 2011; Segura-Puimedon et al., 2014; Stoppel and Anderson, 2017) or Angelman syndrome (Stoppel and Anderson, 2017; Simchi and Kaphzan, 2021), rare neurodevelopmental genetic disorders. Interestingly, these patients show abnormal activity patterns in the OFC. Hypersociability is also found in some mouse models of autism (Yoo et al., 2019). Our data demonstrate that PWSI induced similar behavioral abnormalities in female mice, suggesting that the normal maturation and activity of the OFC_L, at least in females, is important for maintaining an adequate level of sociability in such disorders.

PV expression in the OFC_L is important for female social behavior

Recent studies found important roles of PV expression in the PFC in shaping social and cognitive behaviors. For example, several mouse models of autism that show impaired sociability have shown reduced PV expression in the mPFC (Filice et al., 2016; Kobayashi et al., 2018). Furthermore, maternal separation before weaning (Goodwill et al., 2018) or social isolation during early adolescence (Hinton et al., 2019) reduced PV expression in the OFC and impaired flexibility in behavioral decisions in rodents. Whole-brain knockout of PV caused the mice to show autismlike behavior (Wöhr et al., 2015), but it is still unclear whether the reduced PV expression in subregions of the PFC caused such behavioral deficits. Our study newly applied the CRISPR/Cas9 system to reduce PV expression locally in the OFC_L. Using this method, we confirmed that PV expression in the OFC_L during adolescence is critical for shaping social behaviors in female mice (Fig. 5). This technique can be widely used for studying area-specific PV protein function in various brain areas, although it lacks the reversibility of the loss of function by permanent gene deletion in vivo.

PV is a calcium-binding protein that plays a key role in regulating the physiological properties of PV^+ neurons. In our study,

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decreased the excitability of PV^+ neurons. *I–F* curves measured in the $OFC_1 PV^+$ neurons. sgPV expression caused a significant decrease in current-evoked spike counts in the PV^+ neurons. Two-way ANOVA: F_{AAV} (1,372) = 14.528, ***p < 0.001; $F_{current}$ (11,372) = 121.158, ***p < 0.001; $F_{AAV \times current (11,372)} = 0.388$, p = 0.960. J, Mean half-width of spikes (in ms) measured in PV^+ neurons expressed with the tdTom or the sqPV. Unpaired Student's t test, p = 0.628. **K**, Mean input resistance (in M Ω) measured in PV⁺ neurons expressed with the tdTom or the sgPV. Unpaired Student's t test, p = 0.173. L, Spontaneous EPSCs measured in PV^+ neurons of the OFC₁. Left bars, Mean amplitudes (in pA) of sEPSCs in PV^+ neurons. Unpaired Student's t test, p = 0.636. Right bars, Mean frequency (in Hz) of sEPSCs in PV⁺ neurons. Unpaired Student's t test, p = 0.940. tdTom = 10 cells (5 mice); sgPV = 10 cells (4 mice); female. M, Spontaneous IPSCs measured in excitatory neurons of the OFC₁. Left bars, Mean amplitudes (in pA) of sIPSCs in Pyr neurons. Unpaired Student's t test, **p = 0.006. Right bars, Mean frequencies (in Hz) of sIPSCs in Pyr neurons. Unpaired Student's t test, *p = 0.015. tdTom = 16 cells (3 mice); sgPV = 17 cells (3 mice); female. All data are presented as the mean \pm SEM. For (**D**-**G**): tdTom: n = 6 mice, n = 122 units; sgPV: n = 5mice, n = 61 units. For I-K: tdTom = 16 cells (3 mice); sgPV = 20 cells (4 mice); female. *****p* < 0.001, ***p* < 0.01, **p* < 0.05; n.s., Not significant.



Figure 7. Optogenetic activation of excitatory neurons or inhibition of PV⁺ neurons in the OFC₁ caused hypersociability in female mice. A, Schematic illustration of virus injection and probe implantation for wireless optogenetic manipulation. The virus was injected into the OFC₁ of a mouse. After a week of recovery, the wireless LED probe was implanted in the anterior part of OFC₁ of a mouse. B₁, Schematic illustration of wireless optogenetic manipulation. The smartphone application was used to turn on or off the light by BLE. For optogenetic stimulation, two types of LED light were used at different temporal modes: blue LED (473 nm) at a 20 Hz light-pulse mode; green LED light (532 nm) at a continuous light-on mode. C, Optogenetic activation of excitatory neurons in the female OFG. AAV-CAMKII α -eYFP (eYFP, open bars) or AAV-CAMKII α -ChR2-eYFP (chR2-eYFP, filled bars) was injected into the OFG of PV-Cre female mice. Left, A representative brain image of the OFC, of a WT female mouse injected with the AAV-CAMKII\alpha-ChR2-eYFP. Green, ChR2-eYFP; blue, DAPI. Right, A schematic illustration of a mouse brain section marked with each probe-implanted site in the OFC of ChR2-eYFP-injected female mice. Blue shading, OFC; colored bars, probe-implanted sites confirmed by post hoc histology. **D**, The socialsniffing index of eYFP (open bars) and ChR-2eYFP (filled bars) mice was measured on 2 consecutive days. For Day 1, the light was not delivered (OFF), and the social-sniffing index was not different between the eYFP and ChR2-eYFP groups. Unpaired Student's t test between groups, p = 0.516. For Day 2, the light was delivered to the OFC₁ (ON), and the ChR2-eYFP group showed a higher social-sniffing index than the eYFP group. Unpaired Student's t test between groups, *p = 0.030. Paired Student's t test within a group: eYFP, **p = 0.007; ChR2-eYFP, p = 0.179. E, △Social-sniffing index indicates the difference in social-sniffing index between Day 1 (OFF) and Day 2 (ON). Unpaired Student's *t* test between groups, **p* = 0.036. Wilcoxon signed-rank test for each group: eYFP, *p = 0.036; ChR2-eYFP, p = 0.179. **F**, Locomotion of mice [distance moved (in cm) for 5 min] in the open field chamber was not affected by the optogenetic activation of the OFG. Unpaired Student's t test between groups: OFF, p = 0.267; ON, p = 0.874. Paired Student's t test within a group: OFF versus ON: eYFP, p = 0.078; ChR2-eYFP, p = 0.602. Female: eYFP; n = 5 mice; ChR2, into the OFC₁ of PV-Cre female mice. Left, A representative brain image of the OFC₁ of a PV-Cre female mouse injected with the AAV-EF1*a*-DIO-eArch3.0-eYFP. Green, Arch-eYFP; blue, DAPI. Right, A schematic illustration of a mouse brain section marked with each probe-implanted site in the OFC of Arch-eYFP-injected female mice. Blue shading, OFC; colored bars, probe-implanted sites confirmed by post hoc histology. H, The social-sniffing index of eYFP (open bars) and Arch-eYFP (filled bars) mice. For Day 1, the light was not delivered (OFF), and the social-sniffing index was not different between eYFP and Arch-eYFP groups. Unpaired Student's t test between groups: Day 1, p = 0.950; day 2, #p = 0.062. Paired Student's t test within a group: eYFP, *p = 0.048; ChR2-eYFP, p = 0.480. I, Δ Social-sniffing index of eYFP and Arch-eYFP. Unpaired Student's t test between groups, *p = 0.046. Wilcoxon signed-rank test for each group: eYFP, *p = 0.047; Arch-eYFP, p = 0.547. J, Locomotion of mice [distance moved (in cm) for 5 min] in the open field chamber was not affected by the optogenetic inactivation of the PV⁺ neurons in OFC. Unpaired Student's t test between groups, p = 0.068. Paired Student's t test within a group: OFF versus ON: eYFP, **p = 0.007; Arch-eYFP, **p = 0.003. All data are presented as the mean ± SEM. Dots indicate the individual data. The blue lightning sign indicates the delivery of light stimulation (20 Hz, 473 nm), and the green lightning sign indicates the delivery of light stimulation (continuous, 532 nm). For C-E: eYFP, n = 6; ChR2-eYFP, n = 5; female mice. For G-J: eYFP, n = 7; Arch-eYFP, n = 8; female mice. **p < 0.01, *p < 0.05, #p < 0.1. n.s., Not significant.

knocking down PV decreased the excitability of the PV^+ inhibitory neurons, which in turn decreased synaptic inhibition on nearby excitatory neurons (Fig. 6*H*–*M*). The relative reduction in inhibition compared with excitation in the OFC_L potentially disrupted overall social sniffing-induced activity in the OFC_L (Fig. 6*F*,*G*) and hypersociability (Fig. 5*H*). We further proved this idea by the optogenetic activation of excitatory neurons in the OFC_L that led to hypersociability (Fig. 7). Moreover, a reduced number of PV^+ neurons can also disrupt the network property of the OFC_L , similar to the previous report showing

that the loss of PV^+ neurons correlated with a decrease in the network oscillation in the PFC (Lodge et al., 2009). Maturation of network activity during adolescence may be critical for shaping social sniffing-induced activity in the OFC_L.

In a study of virgin female mice, PV and PNN expression in the auditory cortex was reported to be correlated with learning to retrieve pups (Krishnan et al., 2017); this activity, which occurs in the left auditory cortex, is important for shaping maternal behaviors in female mice (Marlin et al., 2015). Although several studies have shown that the expression of PNN is important for the maturation of PV^+ neurons (Gogolla et al., 2009), we found that expression of the PV but not the PNN is important for the maturation of the female OFC_L during late adolescence. Our data suggest that transcription or translation of PV and PNN may be separately regulated, and adolescent social isolation specifically disrupted PV expression only in a subset of neurons in the OFC_L. Our data further support the idea that plasticity in PV⁺ neurons in the OFC_L is important for shaping female sociability and is also characterized by hemispheric asymmetry. Both genetic and epigenetic mechanisms have been identified as being critical for regulating normal social behavior (Hensch, 2005), and thus social behaviors can be easily disrupted in a wide range of brain psychiatric disorders (Blakemore, 2008). Our study further provides insight into how the disruption of experience-dependent gene expression in the PFC inhibitory neurons, such as reduced PV expression in the OFC_L, may underlie abnormal hypersociability that characterizes certain psychiatric disorders.

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