

1 **Supplementary material for:**

2
3 **Astroglial noradrenaline shapes cerebellar synaptic integration**
4 **and motor adaptation**

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15
16 **Materials and Methods**

17
18 Animal models

19 Experiments were conducted on male and female mice between postnatal day 60 (P60) and P90,
20 in accordance with the European Community Council Directives of 1 January 2013 (2010/63/EU) and
21 with the approval of the local animal welfare committee (certificate B 91 272 108). Animals were housed
22 in a temperature-controlled environment (25 °C) under a 12 h light/dark cycle. All mouse lines were
23 maintained on a C57BL/6J genetic background. When no sex differences were observed, data from male
24 and female mice were pooled for analysis. Several transgenic lines were used in this study. The
25 ALDH1L1-EGFP line expresses the enhanced green fluorescent protein (EGFP) under the control of the
26 astrocyte-specific Aldh1l1 promoter. The aVMAT2cKO mice were kindly provided by Paola Bezzi
27 (University of Lausanne) and generated by crossing the GFAPcre^{ERT2} line expressing a tamoxifen
28 (TAM) inducible cre recombinase transgene driven by the astrocytic glial fibrillary acidic protein
29 (GFAP) promoter with the VMAT2^{lox/lox} line containing cre-excisable loxP sequences in the endogenous
30 VMAT2^{1,2}. Recombination was induced by intraperitoneal injections of tamoxifen (100 mg/kg) for five
31 consecutive days, animals were used after a minimum of 10 days after the first injection. Finally, DBH-
32 cre mice were kindly gifted by Bruno Giros (Paris Cité University) and express the cre recombinase
33 transgene under the dopamine-beta-hydroxylase (DBH) promoter.

34
35 Immunohistochemistry

36 Mice were deeply anesthetized with Dolethal and transcardially perfused with iced-cold
37 buffered saline (PBS) for exsanguination and with a 4% paraformaldehyde in PBS solution for fixation
38 composed of (in mM): 35.4 NaH₂PO₄*2H₂O, 160.7 Na₂HPO₄*2 H₂O, 154 NaCl, pH 7.4. Brains were
39 removed and post-fixed overnight at 4°C, then equilibrated with 30% sucrose at 4°C and frozen for
40 storage at -20°C until immunohistochemistry. Sagittal 35µm thick cerebellar slices were collected using
41 a cryostat. Brain slices were washed with PBS, permeabilized with 0.5% Triton X-100 and labeled with
42 primary antibodies overnight at 4°C. Primary antibodies included: rabbit anti-VMAT2 (1/500, Synaptic
43 Systems), rabbit anti-OCT3 (1/500, Gentauro), mouse anti-NET (1/500, ALOMONE AMT-002), rabbit
44 anti-DAT (1/500, ABCAM AB184451), rabbit anti-DBH (1/500, ABCAM AB209487), mouse anti-
45 calbindin (1/1000; Merck C9848), mouse anti-GS (1/500, Merck MABN543), rabbit anti-TH (1/500,
46 ABCAM AB152), mouse anti-COMT (1/400, BD Biosciences #611970). Following primary antibody
47 incubation, slices were washed twice with PBS before incubation with the appropriate secondary
48 antibody for 2h at room temperature with the following secondary antibodies: goat anti-rabbit 555
49 (1/1000, Thermofischer Scientific #A-21428), goat anti-mouse 488 (1/1000, Thermofischer Scientific

50 #A-11001), goat anti-rabbit 647 (1/1000, Thermofischer Scientific #A-21245). Slices were then washed
51 twice with PBS and once with PB (in mM: 35.4 NaH₂PO₄*2H₂O, 160.7 Na₂HPO₄*2 H₂O) before being
52 mounted in fluoromount (Thermofischer Scientific #18744094). Slices were observed under a confocal
53 Leica SP8 microscope with x10 (0.4 NA), x25 (0.95 NA) and x40 (1.3 NA) objectives. Images were
54 acquired at 12 bit with identical laser power and PMT settings and analyzed with ImageJ and LasX
55 softwares. NE staining (1/500, Immusmol #IS1028) was performed using the STAINperfect
56 Immunostaining kit (ImmuSmol #SP-A-1000, Bordeaux, France), following the manufacturer's
57 protocol: mice were anesthetized and transcardially perfused with a combination of ice-cold 2%
58 paraformaldehyde and the ImmuSmol fixation solution. Coronal and sagittal 35µm slices were collected
59 with a cryostat (free-floating) and immunostaining was performed according to ImmuSmol protocol.
60 Immunostainings were quantified as the proportion of cells identified by Purkinje cell or astrocytic
61 markers that expressed the target protein.

62

63 Magnetic-activated cell sorting (MACS) isolation of astrocytes

64 Mice were deeply anesthetized with isoflurane and euthanized by cervical dislocation. The
65 cerebellar vermis was rapidly dissected for astrocyte isolation and enzymatically dissociated at 37 °C
66 for 30 minutes with the Miltenyi Adult Brain Dissociation Kit (130-107-677, Miltenyi Biotech, France)
67 with gentle trituration every 10 minutes to obtain a single-cell suspension. After filtration through a
68 through a 70 µm SmartStrainer, cells were pelleted by centrifugation (300 × g, 10 minutes, 4 °C) and
69 resuspended in PBS + 0.5 % BSA. Fc receptors were blocked with the accompanying FcR Blocking
70 Reagent, then cells were labeled with Anti-ACSA-2 MicroBeads (130-097-678, Miltenyi Biotech,
71 France) for 15 minutes at 4 °C and loaded onto magnetic selection (MS) columns for positive selection.
72 MS columns were washed twice with PBS + 0.5 % BSA and ACSA-2⁺ astrocytes eluted in 500 µL
73 sorting buffer. For each genotype (WT and aVMAT2cKO), astrocytes isolated from five mice were
74 pooled, counted on an automated cell counter (DeNovix, France), and pelleted by centrifugation (300 x
75 g, 5 minutes).

76

77 Protein quantification by automated capillary immunoassay

78 Cell pellets obtained with the MACS isolation of astrocytes were lysed in 100 µL buffer
79 containing protease and phosphatase inhibitors. Lysates were then incubated on ice for 20 minutes with
80 intermittent mixing and cleared by centrifugation (5000 × g, 15 min, 4 °C). Protein concentrations were
81 determined by Bradford assay (Bio-Rad). Capillary immunoassays were then performed with an
82 “automated western blot system” (WES, ProteinSimple, BioTechne) using the 12-230 kDa Separation
83 Module combined to either the anti-rabbit (#DM-001) or anti-mouse (#DM-00) detection modules
84 depending on the primary antibody. This automated capillary immunodetection delivers antibody-based
85 separation and quantification equivalent to a classical Western blot, with improved throughput and
86 precision. Protein concentration in supernatants of isolated astrocytes was first determined by Bradford
87 assay (Bio-Rad), adjusted to 1 µg/ml in sample buffer provided in the separation module and processed
88 according to the protocol provided by the manufacturer. A six point calibration curve was included to
89 determine the optimal antibody concentration for reliable and quantitative detection. Data analysis was
90 performed using the SimpleWes software, by verifying molecular-weight markers and peak assignment.
91 The area under the curve (AUC) for each peak was exported in excel for subsequent statistical analysis.

92

93 Surgery

94 *Stereotaxic injections of AAVs*

95 Mice anesthesia was induced with 4% isoflurane in 100% O₂, and maintained with 1-1.5%
96 isoflurane. Body temperature was monitored and maintained at 37°C during the whole procedure. The
97 scalp was shaved and sterilized with betadine and 70% alcohol before an incision was made followed
98 by drilling to access the skull. Intracerebral injections were achieved using a glass pipette connected to

99 a syringe pump (Legato 130, KD Scientific). Viruses were delivered at a rate of 0.05-0.1 $\mu\text{L}/\text{min}$. Virus
100 capillary was slowly removed after three minutes post-injection.

101 For the combined expression of the GRAB_{NE2h} sensor and the excitatory DREADD in WT mice,
102 500 nL of AAV9-hSyn-GRAB_{NE2h}-GFP (WZ Biosciences Inc, 2.76×10^{13} GC/ml) and 500 nL AAV5-
103 GFAP-hM3Dq-mCherry (Addgene, #50478, 1.8×10^{13} GC/ml) viruses were injected in lobule VI of the
104 cerebellum (AP: -6.96 mm, ML: 0.0 mm, DV: -0.40 mm). AAV8-Pcp2-hM3Dq-mCherry viral vectors
105 ($> 4.0 \times 10^{13}$ GC/ml) allowing specific excitatory hM3Dq expression in PC were provided by Viral
106 Vector Core (University of Minnesota) and co-injected with the GRAB_{NE2h} sensor mentioned above
107 following the same protocol. DBH-Cre mice were bilaterally injected with 200 nL of AAV9-hSyn-DIO-
108 hM4D(Gi)-mCherry (Addgene, #44362, 2.5×10^{13} GC/ml) viral vectors in the LC (AP: -5.45 mm,
109 ML: = +/-0.85 mm, DV = -3.65 mm) for expression of hM4Di DREADD in NE neurons. CalEx viruses
110 (AAV5-gfaABC1D-mCherry-hPMCA2w/b, Addgene #111568, 8.9×10^{12} GC/ml) were co-injected with
111 GFAP-hM3Dq and either GCaMP6f (AAV5-gfaABC1D-lck-GCaMP6f, Addgene #52924, 2.5×10^{13}
112 GC/ml) or GRAB_{NE2h} in a total volume of 1 μL . All animals were used 3 to 4 weeks after injection.

113

114 *Cranial window and baseplate implantation*

115 For in vivo imaging, a 3 mm diameter circular craniotomy was made above the cerebellar vermis (AP:
116 -6.5 to -7.0 mm; ML: 0.0 mm from Bregma). Three injections of GRAB_{NE2h} virus (300 nL each) in the
117 vermis were performed before the craniotomy was sealed with a sterile glass coverslip (3 mm diameter,
118 #1 thickness), fixed using surgical glue and by dental cement. Mice were allowed to recover for one
119 week. Subsequently, a miniscope-compatible baseplate (Inscopix) was implanted above the cranial
120 window using dental cement to allow miniscope fixation. Mice were then allowed to recover for an
121 additional 2 weeks before starting imaging sessions.

122

123 Preparation of ex vivo cerebellar slices

124 Mice at P60-P80 were deeply anesthetized with isoflurane and sacrificed. The cerebellar vermis
125 was collected after removal of the brain and placed in cold (4°C), oxygenated (95% O₂ + 5% CO₂)
126 cutting artificial cerebrospinal fluid (cutting ACSF, in mM: 110 NaCl, 5 KCl, 2.5 CaCl₂, 1.5 MgSO₄,
127 1.24 KH₂PO₄, 10 D-glucose, 27.4 NaHCO₃, 0.05 D-AP5). The vermis was then sagittally cut at 250 μm
128 using a vibrating blade microtome (650HV, Microm France) for whole-cell patch clamp recordings, and
129 at 400 μm for two-photon imaging. Slices were then transferred to room temperature recording ACSF
130 (in mM: 110 NaCl, 5 KCl, 2.5 CaCl₂, 1.5 MgSO₄, 1.24 KH₂PO₄, 10 D-glucose, 27.4 NaHCO₃) and
131 allowed to recover for 1h prior to recordings.

132

133 Electrophysiological recordings

134 Individual slices were transferred to a submerged chamber mounted on a fixed-stage Scientifica
135 SliceScope Pro 1000 electrophysiology microscope. Slices were continuously perfused with oxygenated
136 ACSF maintained at 30-32°C with a temperature controller managed by the LinLab software
137 (Scientifica), with a flow rate of 2mL/min. Signals were acquired with a MultiClamp 700B amplifier
138 (Molecular Devices) filtered at 1 kHz and recordings were obtained with the Clampex software
139 (Molecular Devices, UK). Spontaneous excitatory and inhibitory postsynaptic currents (sEPSCs and
140 sIPSCs) were recorded from PC by whole-cell patch clamp using borosilicate glass capillaries (Harvard
141 Apparatus, 1.5 OD x 0.86 ID x 100L) with a resistance of 4-5 M Ω , filled with the following intracellular
142 solution in mM: 144 potassium gluconate, 1 MgCl₂, 10 HEPES, 0.5 EGTA (pH 7.4, 280 mOsm). Cells
143 with an access resistance > 20 M Ω at resting potential were excluded from analyses as well as any cell
144 for which a >20 % change in those parameters occurred during the course of the experiment. PC were
145 recorded in voltage-clamp mode with a holding potential of -60 mV. Spontaneous inhibitory post-
146 synaptic currents (sIPSCs) were recorded in the presence of the AMPA receptor antagonist NBQX (10
147 μM , HelloBio #HB0443) and the NMDA receptor antagonist D-AP5 (50 μM , HelloBio #HB0225) while

148 spontaneous excitatory post-synaptic currents (sEPSCs) were recorded in the presence of the GABA_A
149 receptor antagonist gabazine (5 μ M, HelloBio #HB0901). Spontaneous currents were analyzed with
150 Easy Electrophysiology software (Easy Electrophysiology Ltd, UK).

151 Pharmacological block of noradrenergic receptors was obtained by the application of antagonists
152 prazosin (α 1, Merck #P7791-50MG, 30 μ M), propranolol (β , Merck # P0884-1G, 10 μ M), and
153 phentolamine (α , Merck # P7547-100MG, 30 μ M). For experiments with mice expressing DREADD,
154 clozapine N-oxyde (CNO, Sigma-Aldrich #SML2304, 10 μ M) was added to the bath solution.

155 In experiments using mice injected with AAV8-Pcp2-hM3Dq-mCherry, in order to confirm that patch
156 clamped PC also expressed the excitatory DREADD, biocytin (7mg/mL, HelloBio #HB5035) was added
157 to the intrapipette solution and slices were post-fixed overnight in 4% PFA. Slices were then washed
158 and permeabilized before incubation with streptavidin conjugated-488 (ThermoFischer #S32354, 1/500)
159 for 1h45. Post hoc confocal images (Leica SP8) of biocytin-filled PC and virally induced expression of
160 mCherry were acquired to verify their colocalization.

161

162 Two-photon imaging

163 *Acquisition*

164 Ex vivo imaging of cerebellar slices were performed under a two-photon microscope (Femtonics,
165 Budapest) equipped with an 8 kHz resonant scanner combined with a pulse laser (MaiTai-DS,
166 SpectraPhysics, Santa Clara, CA, USA) tuned at 920 nm (green) or 1020 nm (red). Slices were
167 continuously perfused with oxygenated ACSF (in mM: 110 NaCl, 5 KCl, 2.5 CaCl₂, 1.5 MgSO₄, 1.24
168 KH₂PO₄, 10 D-glucose, 27.4 NaHCO₃) and heated at 30-32°C using an automatic temperature controller
169 (TC-324B, Warner Instruments). Images were acquired at 31.1 Hz. Typical imaging protocol consisted
170 in ten minutes of baseline recording followed by fifteen minutes of treatment (e.g. CNO). For evoked
171 NE-release, a stimulating electrode (borosilicate glass capillary from Phymep (1.5 OD x 0.86 ID x 100L)
172 2-3 M Ω) filled with ACSF was positioned in the molecular layer of the cerebellar cortex. Stimulation
173 was delivered using a DS2A voltage stimulator - Mk.II (Digitimer) under software control, as a single
174 train of 20 Hz. One stimulation was acquired prior to any treatment (baseline) and CNO was bath applied
175 for 15 min before another train of stimulation without changing the position of the stimulating electrode.
176 Calibration-curve protocols were acquired as followed: NE or DA was applied in increasing
177 concentrations (in μ M): 0, 0.01, 0.1, 0.25, 0.5, 1, for 3 min per concentration.

178

179 *Analysis*

180 Videos were exported in AVI format and averaged at 9 frames per second to reduce noise and
181 improve signal quality. Analysis was performed using a custom-built Python pipeline called DETECT
182 (Dynamic Extraction and Tracking of Emitted Cellular Transients)³, specifically designed for an
183 automatic, GMM-based (Gaussian Mixture Model) detection of spontaneous NE or GCaMP signals
184 depending on the experiment. This pipeline identifies individual fluorescence clusters while correcting
185 for background noise and motion correction. Following detection, we used TrackMate in Fiji (ImageJ)
186 to extract a range of spatiotemporal features from each identified cluster, including area, perimeter, mean
187 intensity, total number of events, and directionality of propagation. The resulting csv files were further
188 processed and analyzed in MATLAB using custom-written scripts for group-level statistical
189 comparisons and visualization. For evoked NE-release, videos were exported and treated in the same
190 way, and stimulation response area and intensity was quantified with a custom-built matlab code.
191 Calibration-curve data were extracted as above and analyzed with a custom-built python code where
192 fluorescence movies were processed frame-by-frame. For each frame, the mean fluorescence of non-
193 zero pixels was calculated to exclude static background, and $\Delta F/F$ was computed using the first 60 s of
194 the drug-free period as F_0 to minimize bleaching effects. Traces were **smoothed** with a centered moving
195 average. Pharmacological responses were quantified by fitting $\Delta F/F$ time courses with a four-parameter
196 logistic model using nonlinear least-squares regression. Fits typically achieved R² values of ~0.92, and

197 parameter stability was confirmed by sensitivity analyses with 5% added noise. Baseline values were
198 estimated over a 180 seconds pre-stimulation period.

199 Behavioral experiments

200 *Treadmill-based locomotion assessment*

201 To assess forced locomotion, WT and aVMAT2cKO mice were tested on a 5-lane treadmill
202 (#8710RTS Panlab Harvard Apparatus). Both groups, WT and aVMAT2cKO mice, underwent the
203 tamoxifen injections procedure. Each treadmill lane measured 532 mm in length, 100 mm in width and
204 50 mm in height. The treadmill was equipped with a control unit (LE8700TS Treadmill Control Unit,
205 Panlab Harvard Apparatus) that allowed precise adjustment of speed parameters. The protocol, adapted
206 from ⁴, starts with a stationary period (0 cm/s) for 1 minute, followed by an initial acceleration to 10
207 cm/s over 30 seconds. Mice were then maintained at 10 cm/s for three intervals: from 1.5 to 2 minutes,
208 3 to 4 minutes, and 5 to 6 minutes, each separated by 1 minute rest phases (0 cm/s). At 7 minutes, the
209 speed was increased to 20 cm/s for 1 minute. The total duration of the protocol was 8 minutes.

210

211 *Horizontal ladder test*

212 To assess skilled locomotion and sensorimotor coordination, WT and aVMAT2cKO mice were
213 subjected to the ladder rung walking test as previously described ⁵. Both groups, WT and aVMAT2cKO
214 mice, underwent the tamoxifen injections procedure. The apparatus consisted of two transparent
215 plexiglass side walls (100 cm long, 20 cm high) and evenly spaced metal rungs (1 cm apart) forming a
216 ladder elevated approximately 1 meter above the ground. Irregular sessions consisted of irregular
217 spacing by randomly removing rungs in order to prevent pattern learning and challenge motor
218 coordination. Each regular and irregular session consisted of 3 trials each. For irregular sessions, 3
219 random patterns were created and maintained identical for each mouse. Mice were familiarized to the
220 task by performing 3 regular sessions prior to the tests. The following day, mice were tested with 3
221 regular sessions and 3 irregular sessions. They were placed at one end of the ladder and allowed to
222 traverse to their home cage located at the opposite end. Crossings were recorded using a high-speed
223 camera (200 fps, acA2040-120um, Basler Inc.) integrated into a custom closed-loop system that
224 automatically adjusted the camera position to precisely track the mice as they traversed the ladder. The
225 camera was positioned laterally and slightly inclined to capture precise paw movements. Videos were
226 analyzed frame-by-frame (MPC software) to evaluate forelimb and hindlimb placements using a 7-
227 points foot fault scoring system ⁵:

- 228 - 0: correct placement
- 229 - 1: partial placement, paw is misplaced with only toes or heels
- 230 - 2: correction, paw aims for one rung but is placed on another without touching the first
- 231 - 3: replacement, paw position is adjusted after first initial placement
- 232 - 4: slight slip without fall
- 233 - 5: deep slip, paw slips leading to imbalance and fall
- 234 - 6: total miss, paw misses the rung leading to a fall.

235 For analysis, errors were expressed as a percentage of total steps to account for potential differences in
236 step count across trials. The total number of errors per session and the corresponding error score intensity
237 were measured.

238

239 GRAB_{NE2h} miniscope imaging

240 To control for potential bias from the surgery or miniscope attachment on motor performances, mice
241 performed the treadmill and horizontal ladder protocol before and after surgery for GRAB_{NE2h} imaging.
242 Imaging acquisition during the task was performed using the nVista miniscope system (Inscopix), fixed
243 on the surgically implanted baseplate, with a 20 frames per second acquisition rate. Raw signals were
244 then processed in Inscopix Data Processing Software as followed: after motion correction, regions of

245 interest (ROIs) were manually defined based on signal quality while avoiding blood vessels, using the
246 same ROI size across videos. $\Delta F/F$ traces were then extracted and normalized for each ROI by their root
247 mean square, and aligned to the behavioral videos for correlation analyses. Correlation analyses, power
248 spectral density estimates, and statistical tests were performed in MATLAB.

249

250 Statistics

251 All data were analyzed using GraphPad Prism, MATLAB, and Python. Data normality was assessed
252 using the Shapiro-Wilk test. Outliers were identified using the Robust Regression and Outlier Removal
253 (ROUT) method ($Q = 1\%$)⁶ implemented in GraphPad Prism, which controls the false discovery rate,
254 and were removed prior to statistical analysis. For comparisons between two groups, two-tailed paired
255 or unpaired Student's t tests were used when data were normally distributed. Otherwise, the Wilcoxon
256 signed-rank test (paired) or Mann-Whitney U test (unpaired) was applied. One- or two-way ANOVA,
257 including repeated-measures designs where appropriate, were used, followed by post hoc multiple-
258 comparison tests using Tukey's or Sidak's methods as appropriate. All data are reported as mean \pm SEM.
259 Statistical significance was set at $\alpha = 0.05$. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$ and **** $p < 0.0001$.

260

261 **References**

262

- 263 1. Petrelli, F. *et al.* Disruption of Astrocyte-Dependent Dopamine Control in the Developing Medial
264 Prefrontal Cortex Leads to Excessive Grooming in Mice. *Biol. Psychiatry* **93**, 966–975 (2023).
- 265 2. Petrelli, F. *et al.* Dysfunction of homeostatic control of dopamine by astrocytes in the developing
266 prefrontal cortex leads to cognitive impairments. *Mol. Psychiatry* **25**, (2020).
- 267 3. Niu, W. *et al.* Sharp and Fast Dynamic Extraction and Tracking of Emitted Cellular Transients. *bioRxiv*
268 2026.04.16.718018 (2026) doi:10.64898/2026.04.16.718018.
- 269 4. Paukert, M. *et al.* Norepinephrine controls astroglial responsiveness to local circuit activity. *Neuron* **82**,
270 1263–1270 (2014).
- 271 5. Metz, G. A. & Wishaw, I. Q. The ladder rung walking task: a scoring system and its practical
272 application. *J. Vis. Exp.* <https://doi.org/10.3791/1204> (2009) doi:10.3791/1204.
- 273 6. Motulsky, H. J. & Brown, R. E. Detecting outliers when fitting data with nonlinear regression - a new
274 method based on robust nonlinear regression and the false discovery rate. *BMC Bioinformatics* **7**, (2006).

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Extended data

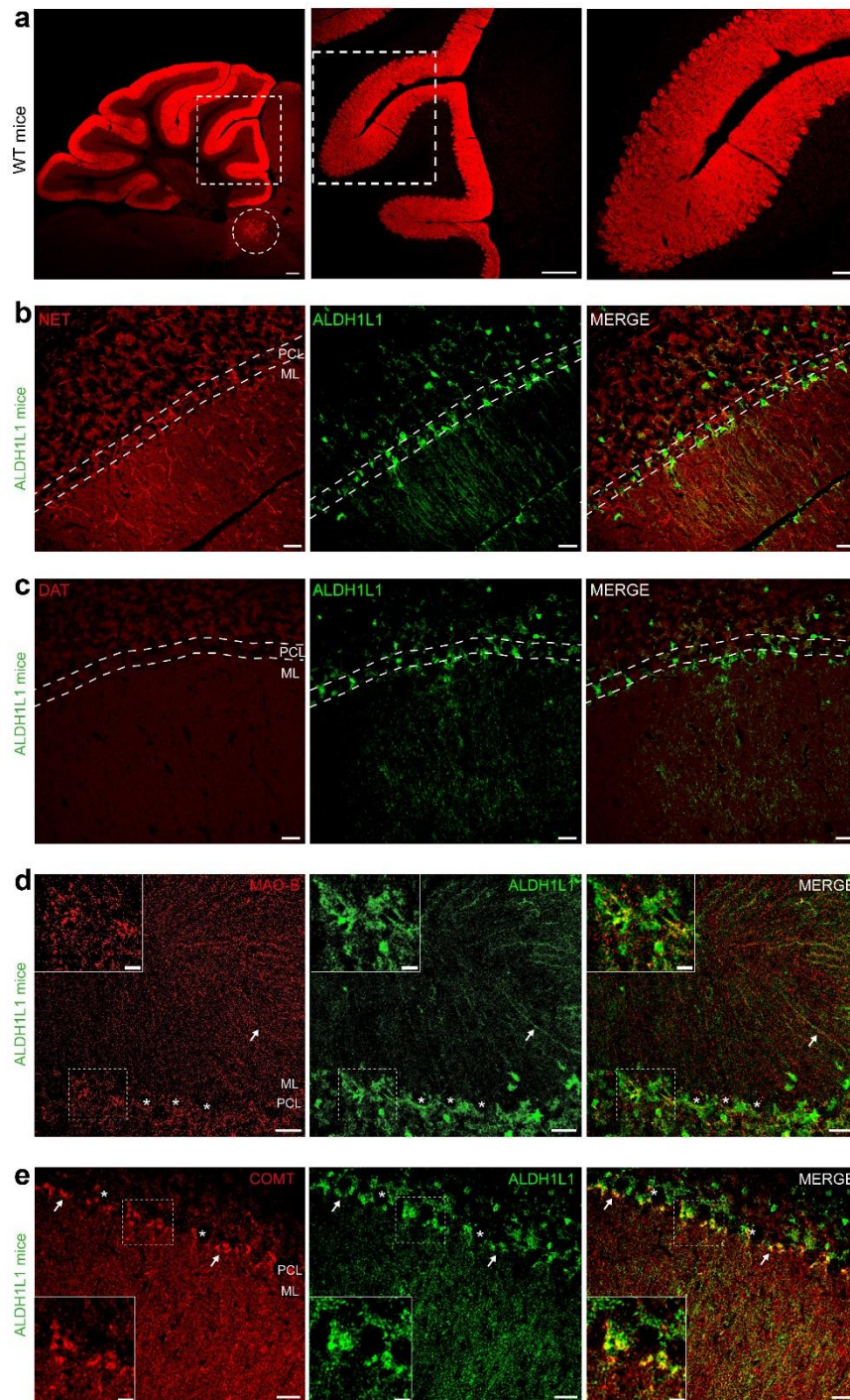


Figure 1. Monoaminergic signaling components in the cerebellar cortex. **a.** Tyrosine hydroxylase (TH, red) immunostaining in WT mice showing widespread labeling across cerebellar lobules. Dashed squares indicate regions shown at higher magnification (middle and right panels). The dashed circle highlights TH-positive *locus coeruleus* neurons (n = 8). Scale bar, 100 μ m. **b.** NET immunostaining (red) in ALDH1L1-eGFP mice (green) showing NET expression restricted to sparse fibers within the molecular layer and absent from Bergmann glia (BG). Scale bar, 25 μ m (n=4). Dashed lines delineate the Purkinje cell layer (PCL) and molecular layer (ML). **c.** DAT immunostaining (red) in ALDH1L1-eGFP mice (green) showing absence of DAT expression in the cerebellar cortex. Scale bar, 25 μ m (n=2). Dashed lines delineate the Purkinje cell layer (PCL) and molecular layer (ML). **d.** MAO-B immunostaining (red) in ALDH1L1-eGFP mice (green) showing expression of monoamine degradation enzymes in BG. Insets show higher magnification views. Scale bars, 25 μ m (main), 10 μ m (inset) (n=2). **e.** COMT immunostaining (red) in ALDH1L1-eGFP mice (green) confirming expression of monoamine degradation pathways in BG. Insets show higher magnification views. Scale bars, 25 μ m (main), 10 μ m (inset) (n=3).

422 **Supplementary video captions**

423

424 **Video 1. Detection of spontaneous NE transients in the cerebellar cortex.** Left: Raw two-photon
425 GRAB_{NE2h} imaging showing spontaneous NE transients. Middle: Background removal using our custom
426 pipeline (DETECT) (10), isolating spontaneous NE transients. Right: Detection and segmentation of NE
427 transients for quantitative analysis. Scale bar, 25 μ m. Displayed at 20x real-time speed.

428

429 **Video 2. Extracellular Ca²⁺ removal abolishes spontaneous NE release.** Left: Two-photon GRAB_{NE2h}
430 imaging of spontaneous NE transients under baseline conditions. Right: Two-photon imaging of
431 spontaneous NE transients after extracellular Ca²⁺ removal. Scale bar, 25 μ m. Displayed at 20x real-
432 time speed.

433

434 **Video 3. Chemogenetic silencing of LC noradrenergic neurons suppresses evoked NE transients**
435 **in the cerebellar cortex.** Left: Two-photon GRAB_{NE2h} imaging of electrically evoked NE release in
436 the molecular layer under baseline conditions. Right: Electrical stimulation after hM4Di chemogenetic
437 silencing of LC noradrenergic neurons with CNO (10 μ M, 30 min), showing suppression of evoked NE
438 transients. Scale bar, 25 μ m. Displayed at 20x real-time speed.

439

440 **Video 4. Chemogenetic silencing of LC noradrenergic neurons does not abolish spontaneous NE**
441 **release.** Left: Two-photon GRAB_{NE2h} imaging of spontaneous NE release under baseline ACSF
442 conditions. Right: Two-photon GRAB_{NE2h} imaging following chemogenetic silencing of LC
443 noradrenergic neurons with CNO (10 μ M, 30 min), showing persistent spontaneous NE release. Scale
444 bar, 25 μ m. Displayed at 20x real-time speed.

445

446 **Video 5. Chemogenetic activation of Bergmann glia increases spontaneous NE release.** Left: Two-
447 photon GRAB_{NE2h} imaging of spontaneous NE transients under baseline ACSF conditions. Right: Two-
448 photon GRAB_{NE2h} imaging during hM3Dq chemogenetic activation of Bergmann glia with CNO (10
449 μ M), showing increased spontaneous NE transients. Scale bar, 25 μ m. Displayed at 20x real-time speed.

450

451 **Video 6. Chemogenetic activation increases astroglial Ca²⁺ dynamics.** Left: Two-photon GCaMP6f
452 imaging of Bergmann glia Ca²⁺ transients under baseline ACSF conditions. Right: Two-photon
453 GCaMP6f imaging during astroglial hM3Dq chemogenetic activation with CNO (10 μ M), showing
454 increased astroglial Ca²⁺ dynamics. Scale bar, 25 μ m. Displayed at 5x real-time speed.

455

456 **Video 7. Disruption of astroglial Ca²⁺ signaling reduces spontaneous NE transients and suppresses**
457 **hM3Dq-induced increases in spontaneous NE release.** Left: Two-photon GRAB_{NE2h} imaging of
458 spontaneous NE release under baseline ACSF conditions in mice expressing the calcium extruder CalEx
459 in astroglial cells. Right: Two-photon GRAB_{NE2h} imaging during chemogenetic activation of Bergmann
460 glia with CNO (10 μ M) in mice expressing CalEx in astroglial cells, showing suppression of the hM3Dq-
461 induced increase in spontaneous NE release. Scale bar, 25 μ m. Displayed at 20x real-time speed.

462

463 **Video 8. CalEx suppresses astroglial Ca²⁺ dynamics during chemogenetic activation.** Left: Two-
464 photon GCaMP6f imaging of astroglial Ca²⁺ transients under baseline ACSF conditions in mice
465 expressing the calcium extruder CalEx in cerebellar cortex astrocytes. Right: Two-photon GCaMP6f
466 imaging during chemogenetic activation of Bergmann glia with CNO (10 μ M) in mice expressing CalEx
467 in astrocytes, showing reduced Bergmann glia Ca²⁺ dynamics. Scale bar, 25 μ m. Displayed at 5x real-
468 time speed.

469

470 **Video 9. aVMAT2cKO mice show impaired motor coordination during the horizontal ladder task.**
471 Top: WT mouse performing the horizontal ladder task under irregular conditions. Bottom:
472 aVMAT2cKO mouse performing the same task, showing increased fine and strong stepping errors.
473 Displayed at 1x real-time speed.

474