

Supplementary information

Environmental enrichment induces behavioural disturbances in neuropeptide Y knockout mice

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1 **Supplementary Materials and Methods**

2 **EPM.** For the EPM mice were placed in the centre of a maze consisting of a central platform (5 x 5
3 cm), two opposing open arms (30 cm long and 5 cm wide), and two opposing closed arms of the
4 same size but equipped with 15-cm high walls at their sides and the far end¹. At the beginning of
5 each trial, the animals were placed on the central platform facing an open arm. The behaviour of the
6 animals during a 5-min test period was videotaped and tracked with VideoMot 2 software (TSE
7 Systems, Bad Homburg, Germany). Number of entries into and time spent on the open and closed
8 arms as well as total distance travelled was quantified to evaluate anxiety, exploration and
9 locomotion.

10 **OF.** The OF consisted of a box (50 x 50 x 30 cm) made of opaque grey plastic. The ground area of the
11 box was divided into a 36 x 36 cm central area and the surrounding border zone². Mice were
12 individually placed in the centre of the OF, and their behaviour during a 5-min test period was
13 videotaped and tracked with the VideoMot 2 software. Number of entries into and time spent in the
14 central area were used as anxiety readouts, while total distance travelled was measured to analyse
15 locomotion.

16 **Stress-induced hyperthermia.** This test consisted of a series of 2 rectal temperature measurements³.
17 For this, the basal rectal temperature (T1) was measured with a digital thermometer (BAT-12,
18 Physitemp Instruments, Clifton, New Jersey, USA) equipped with a rectal probe for mice representing
19 a physical stressor. Fifteen min later a second measurement of the temperature (T2) was performed.
20 T2 was regarded as a measure of stress reactivity.

21 **FST.** For the FST we used a glass beaker (inner diameter 18 cm, height 27 cm) containing tap water at
22 25 °C². The water was 20 cm deep, which prevented the mice from touching the bottom of the
23 beaker. Mice were carefully put into the water-filled beaker for 6 min to measure the time of
24 immobility (passive floating in the water), swimming and climbing. For this, each session was

1 videotaped and manually scored by a trained investigator blind to the treatment. Time of immobility
2 was used to evaluate depression-like behaviour.

3 **BM.** The BM consisted of a 0.8 cm thick white PVC circular board (diameter: 91 cm) with 20 equally-
4 spaced holes (diameter: 5 cm) at a distance of 2.5 cm from the edge. This apparatus was positioned
5 61.5 cm above floor level and a black escape box (11 x 5 x 5 cm), to which the mice gained access by
6 stepping down through the target hole, could be fitted below any of the holes. To evaluate learning
7 and memory a slightly modified version of the protocol provided by Attar et al.⁴ was used. Briefly, on
8 day 1 of a 2-week test period mice were habituated to the maze by placing them on the board under
9 a clear glass beaker. Mice under the beaker were then carefully guided to the target hole, which
10 leads into the escape box, and allowed to enter on their own. If the mice did not enter, the way to
11 enter was shown to them by the investigator. To avoid potential odour cues the maze and box were
12 thoroughly cleaned after each trial and the position of the target box was changed as well. On the
13 next 2 days (training phase) mice were allowed to freely explore the maze in 3 (day 2) and 2 (day 3)
14 2-min trials. To assess learning capabilities we measured the latency to find the target hole and the
15 number of errors before identifying the target hole (primary errors). If the mice did not find the
16 target hole, they were again carefully guided to the target hole under the glass beaker. The latency to
17 approach the first hole was used to identify any influence of anxiety-like behaviour. Four different
18 visual cues were mounted on each wall of the testing room to facilitate learning. Fortyeight h after
19 the last training trial the hidden escape box was removed and short-term memory was estimated by
20 the time spent in the target quadrant (tracked with VideoMot 2 software). Seven days after the first
21 probe trial, a second probe trial was performed to evaluate long-term memory.

22 **Cage behaviour.** For this test, mice were put individually into a polycarbonate cage of size IV
23 measuring 59.0 x 38.0 x 20.0 cm (length x width x height) and video recorded for 10min. The cage
24 contained 5 items: a running wheel (Dehner, Graz, Austria), a wooden tunnel (Dehner)⁵, a hay tunnel
25 (Dehner) and two transparent red plastic mouse houses (Ehret, Tulln, Austria). The time each object
26 was used by the mice was quantified.

1 **CORT ELISA.** Circulating CORT levels were determined with an enzyme immunoassay kit (Assay
2 Designs, Ann Arbor, Michigan, USA). For this, mice were euthanized with an overdose of
3 pentobarbital (150 mg/kg IP) and blood samples were collected by cardiac puncture with syringes
4 containing sodium citrate (3.8 %; w/v) as an anticoagulant. Following centrifugation for 10 min at 4
5 °C and 7000 rpm, blood plasma was stored at -20 °C until assay. The assay was performed according
6 to the manufacturer's instructions. According to the manufacturer's specifications, the sensitivity of
7 the assay is 27 pg/ml, and the intra- and inter-assay coefficient of variation amounts to 7.7 and 9.7 %,
8 respectively.

9 **Mouse brain microdissection.** For peptide, protein and mRNA measurements brains were
10 microdissected as previously described⁶. Briefly, fresh frozen mouse brains were cut manually into
11 approximately 1 mm thick coronal slices from which brain areas of interest were isolated on a cold
12 plate (Weinkauf Medizintechnik, Forchheim, Germany) set at -20 °C under a stereomicroscope.
13 Hippocampal tissue was collected across the whole rostrocaudal extent starting from the limit of the
14 hippocampal formation (Bregma: -0.94) to the caudal end of the dentate gyrus (Bregma: -4.04). The
15 microdissection of the amygdala was also performed across the whole rostrocaudal extent of the
16 amygdalar complex (Bregma: -0.58 to Bregma: -2.54) and thus the tissue taken contained all major
17 amygdalar subnuclei (medial, central, basolateral, basomedial, lateral and cortical). Isolated brain
18 areas were collected in homogenization tubes kept on dry ice and stored at -70 °C until further
19 processing. Areas isolated from the left and right hemisphere were collected in separate tubes. One
20 brain half was used for RNA extraction, while the other half was used for protein extraction. To
21 exclude lateralization effects, a random number generator was used to allocate the right or the left
22 brain half to RNA or protein extraction, respectively.

23 **Protein extraction.** Brain areas used for protein and peptide measurements were homogenized in
24 lysis buffer (50 mM Tris-HCl pH 8, 150 mM NaCl, 1 % (v/v) Triton X-100, 0.5 % (v/v) sodium
25 deoxycholate and 10 mM PMSF) using a Peqlab Precellys 24 homogenizer (Peqlab, Erlangen,
26 Germany)⁷. The tissue homogenates were centrifuged (10,000 rpm, 4 °C, 10 min) to pellet debris and

1 the protein content was measured with the BCA protein assay kit (Pierce Biotechnology, Rockford, IL,
2 USA).

3 **Western Blot.** For Western blot, protein lysates (10 or 50 µg; amygdala or hippocampus) were mixed
4 with 4x NuPAGE LDS sample buffer and NuPAGE sample reducing agent, heated for 10 min at 70 °C
5 and loaded on 4-12 % gradient SDS-PAGE gels (Life Technologies, Carlsbad, CA, USA) for
6 electrophoresis⁷. After transfer to nitrocellulose membranes (45 µm; Life Technologies), the
7 membranes were blocked with non-fat milk (5 % w/v) in Tris-buffered saline containing Tween20
8 (TBS) for 1 h. After being washed, membranes were incubated overnight at 4°C with anti-NR3C1
9 antibody (1:1000, Cell Signalling, D6H2L) diluted in TBS containing bovine serum albumin (5 % w/v).
10 After another wash, the membranes were incubated with horseradish peroxidase-conjugated goat
11 anti rabbit IgG (1:200,000, Biomol). Immunoreactive bands were visualized with Clarity™ Western
12 ECL Blotting Substrate (Biorad, Hercules, CA, USA) and detected by the ChemiDoc™ Touch Imaging
13 System (Bio-Rad). For normalization, membranes were incubated with anti-glyceraldehyde 3-
14 phosphate dehydrogenase (GAPDH) antibody (1:1000, Santa Cruz, SC-25778). Densitometric
15 evaluations of immunoreactive bands were performed with the Image Lab software volume module
16 (Biorad).

17 **BDNF ELISA.** Amygdalar and hippocampal BDNF concentrations were measured using the BDNF E_{max}[®]
18 ImmunoAssay System (Promega, Madison, WI, USA). The assay was performed according to the
19 manufacturer's instructions using protein lysates diluted in assay buffer (1:10). The company reports
20 a sensitivity of 15.6 pg/ml and a cross reactivity with other related neurotrophic factors of less than 3
21 % for this assay.

22 **NPY fluorescence EIA.** NPY concentrations were measured by fluorescence EIA (Phoenix
23 Pharmaceuticals, Burlingame, CA, USA). For this, NPY was extracted from the amygdalar and
24 hippocampal protein lysates. Protein (200 or 500 µg) was extracted in 2N acetic acid, respectively,
25 and centrifuged for 10 min at 2400 rpm/ 4 °C⁸. The supernatants were lyophilized and stored at -70

1 °C. For the assay the lyophilisate was reconstituted in assay buffer and the EIA was performed
2 according to the manufacturer's instructions. The sensitivity of the assay is 11.9 pg/ml and there is no
3 cross reactivity with peptide YY or pancreatic polypeptide.

4 **Real-time RT PCR.** Brain halves used for real-time RT PCR were homogenized in QIAzol lysis reagent.

5 RNA was extracted with the RNeasy Lipid Tissue Mini kit (Qiagen, Hilden, Germany) and reverse
6 transcribed using the high capacity cDNA reverse transcription kit (Applied Biosystems, FosterCity,
7 CA, USA)⁶. For relative quantification of mRNA levels within a given brain area, specific primers for
8 *Npy* (forward 5' -> 3': CAGATACTACTCCGCTCTGCGACTACA; reverse 5' -> 3':

9 TTCCTTCATTAAGAGGTCTGAAATCAGTGTC), *Bdnf* (forward 5' -> 3': GTGACAGTATTAGCGAGTGG;

10 reverse 5' -> 3': TTCTCTAGGACTGTGACCGT), *Nr3c1* (forward 5' -> 3': GACTCCAAAGAATCCTTAGCTCC;

11 reverse 5' -> 3': CTCCACCCCTCAGGGTTTTAT), *Gapdh* (Qiagen, Mm_Gapdh_3_SG

12 QuantiTectPrimerAssay), phosphoglycerate kinase 1 (*Pgk1*; forward 5' -> 3':

13 ATGTCGCTTTCCAACAAGCTG; reverse 5' -> 3': GCTCCATTGTCCAAGCAGAAT) and peptidylprolyl

14 isomerase A (*Ppia*; forward 5' -> 3': TTCCAGGATTCATGTGCCAG; reverse 5' -> 3':

15 CCATCCAGCCATTCAGTCTT) were used at a concentration of 1 µM each together with the

16 SsoAdvanced Universal SYBR Green Supermix (Bio-Rad). DNA sequencing and BLAST analysis

17 confirmed the amplification of the correct base pair sequence. Samples were measured in triplicates

18 on a LightCycler 480 System (Roche, Rotkreuz, Switzerland). Quantitative measurement of target

19 gene levels relative to controls was performed with the $2^{-\Delta\Delta Ct}$ method. *Gapdh*, *Pgk1* and *Ppia* were

20 used as reference genes.

21 **Electron microscopy.** Tissue preparation for electron microscopy followed our previously published

22 procedure⁹. In brief, excised mouse brains were cut in half at the midline, and immersion-fixed for 48

23 h in 2 % formaldehyde, 2.5 % glutardialdehyde in 0.1 M cacodylate buffer pH 7.4 at 4 °C and rinsed in

24 the same buffer for at least 24 h. Coronal vibratome sections were made on a Leica VT 1000

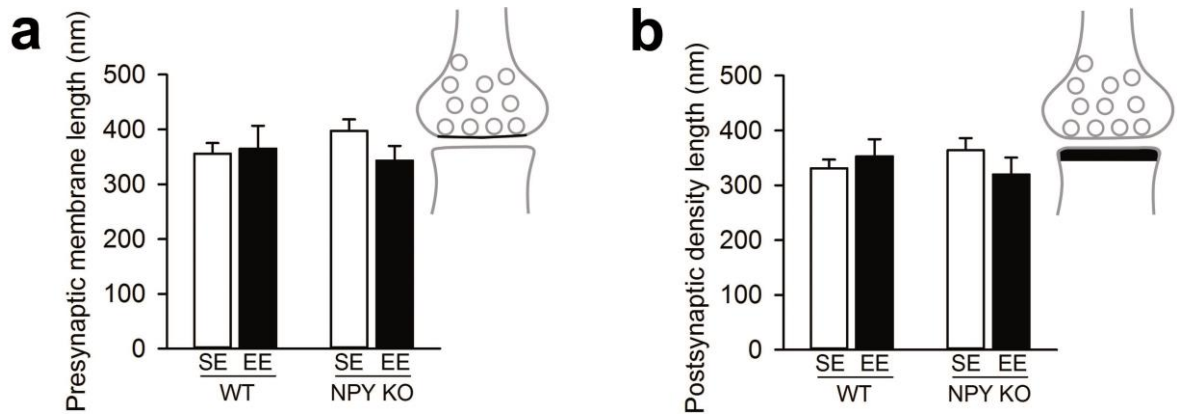
25 vibratome until the DGpl was reached, verified by comparing with thionine-stained vibratome

26 sections and the mouse brain atlas of Franklin and Paxinos¹⁰. A vibratome section close to Bregma -

1 1.3 was cut at 150 μm thickness, post fixed in 2 % osmium tetroxide solution, dehydrated in a
2 series of graded alcohols, and embedded in TAAB embedding resin (TAAB, Berkshire, UK). Three
3 coherent semi thin sections were made at a thickness of 500 nm and stained with 1 % toluidine blue
4 solution, of which the first and third were used for counting the number of cells in the DGpl.
5 Immediately adjacent to these semi thin sections, serial thin sections were made at 55 nm thickness,
6 of which one pair was used for further analysis. These thin sections were examined with a Tecnai G2
7 20 electron microscope (FEI, Eindhoven, Netherlands) operating at 120 kV with a US 1000 digital
8 camera (Gatan, Pleasanton, USA). For analysis, a minimum of 20 image pairs were made from two
9 adjacent sections in each animal, using several software packages as previously described⁹. Serial EM
10 software (version 3.3.1) was used to obtain and merge 4 adjacent camera images at 5000x
11 magnification. FEI Serial Section Software (FEI, Eindhoven Netherlands) was used to store the
12 location of each point on the first image of each pair and to retrieve its location on the second image,
13 where another montage was obtained using Serial EM. The numerical densities of synapses and DCV
14 were assessed using a dissector of 5.5 x 5.5 μm size. Furthermore, on 10 synapses/animal, the
15 following synaptic features were determined on single sections: mean length of the presynaptic
16 membrane cross section and of the postsynaptic density in cross section, average width of the
17 synaptic cleft as well as the average number of docked vesicles (vesicles with a maximum distance
18 from the presynaptic membrane of one vesicle diameter) and undocked vesicles (those with a
19 maximum distance of one vesicle diameter from docked or other undocked vesicles at the same
20 synapse) within the thin section. The ObjectJ platform of ImageJ was used to measure these
21 parameters. All measurements and counts were made on coded samples by an experimenter blind to
22 the treatment groups.

1 **Supplementary Figure**

2



4 **Fig. S1. EE and NPY KO do not alter (A) presynaptic membrane length and (B) postsynaptic density**
5 **length of synapses in the DGpl.** Values were measured in WT and NPY KO mice kept in EE or SE and
6 represent mean \pm SEM. Data in A and B are derived from 10 synapses/animal and 4-5 animals/group.

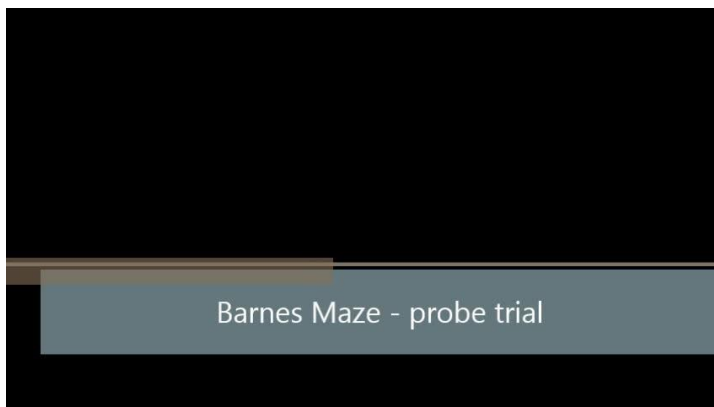
7

1 **Supplementary Movies**



2

3 **Movie S1. Representative behaviour of WT and NPY KO mice during the first training trial of the**
4 **Barnes Maze test.** *NPY KO mice move very slowly and tend to freeze in the centre of the maze. In*
5 *contrast, WT mice immediately begin to explore the maze after starting the training trial.*



6

7 **Movie S2. Representative behaviour of EE-housed and SE-housed NPY KO mice during the probe**
8 **trial of the Barnes maze test.** *EE-housed NPY KO mice do not remember the location of the target*
9 *hole and thus employ a random search strategy. SE-housed NPY KO mice remember the location of*
10 *the target hole and stay in the target quadrant.*

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