# **Expanded View Figures**



GAPDH 37 kDa

Figure EV1.

# Figure EV1. Mecp2 levels and axonal vesicle number in cortical Mecp2-silenced neurons.

- A Western blot analysis of DIV 5 cortical neurons transfected with siMecp2 or siControl (siCtl) to assess Mecp2 protein levels.
- B We quantified the number of BDNF-mCherry-containing vesicles trafficking within cortical axons transfected with siMecp2 or siControl (siCtl). Data are presented as the mean  $\pm$  SEM of at least three independent experiments, with 94 siCtl and 81 siMecp2 axons and 894 siCtl and 753 siMecp2 vesicles (unpaired *t*-test).
- C Brain extracts from WT, WT HTT<sub>SD</sub>, and WT HTT<sub>SA</sub> mice were analyzed by Western blotting for total HTT and phospho-HTT. Quantification of n = 3 samples per genotype did not reveal any significant differences on total HTT. The mutation of HTT S421 into alanine or aspartic acid abrogated the recognition of phospho-HTT by the phospho-specific antibody, further demonstrating the specificity of HTT phosphorylation at S421. Data are presented as means  $\pm$  SEM, and one-way ANOVA test followed by Tukey's post-hoc analysis for multiple comparisons.
- D We quantified the number of BDNF-mCherry-containing vesicles trafficking within WT,  $\text{HTT}_{SA}$ , or  $\text{HTT}_{SD}$  cortical axons. Data are presented as the mean  $\pm$  SEM of at least three independent experiments, with 80 axons and 700 vesicles per condition (one-way ANOVA test followed by Tukey's post-hoc analysis for multiple comparisons).
- E Western blot analysis of DIV 5 HTT<sub>SA</sub> or HTT<sub>SD</sub> cortical neurons transfected with siMecp2 or siControl (siCtl) to assess Mecp2 protein levels.
- F Quantification of the number of BDNF-mCherry-containing vesicles trafficking within WT, HTT<sub>SA</sub>, or HTT<sub>SD</sub> cortical axons transfected with siMecp2 or siControl (siCtl). Data are presented as the mean  $\pm$  SEM of at least three independent experiments, with WT (n = 753 vesicles/81 axons), HTT<sub>SA</sub> (n = 812 vesicles/81 axons), or HTT<sub>SD</sub> (n = 787 vesicles/82 axons) (one-way ANOVA test followed by Tukey's post-hoc analysis for multiple comparisons). ns = not significant.
- G Representative Western blot analysis of PSD-95 and phospho-TrkB protein levels in striatum samples from WT, KO HTT<sub>SD</sub>, KO WT, and KO HTT<sub>SA</sub> mice.

Source data are available online for this figure.

#### Figure EV2. Phenotypic characterization of WT, HTT<sub>SD</sub>, HTT<sub>SA</sub> (A, B, C, and D) mice and WT, KO/HTT<sub>SD</sub>, KO/HTT<sub>SA</sub> (E, F, G, and H) mice.

- A There were no significant differences in body weight between 6-month-old WT, HTT<sub>SD</sub>, and HTT<sub>SA</sub> mice (WT: n = 11; HTT<sub>SD</sub>: n = 6, and HTT<sub>SA</sub>: n = 10).
- B No significant differences in the distance travelled in the open-field between 4-month-old WT, HTTSD, and HTTSA mice (WT: n = 10; HTT<sub>SD</sub>: n = 6 HTT<sub>SA</sub>: n = 11).
- C No differences between genotypes in the accelerating Rotarod test for assessing motor coordination of 6-month-old WT (n = 11), HTT<sub>SD</sub> (n = 11), and HTT<sub>SA</sub> mice (n = 12).
- D No differences in forelimb strength of 6-month-old WT (n = 11), HTT<sub>SD</sub> (n = 11), and HTT<sub>SA</sub> mice (n = 12) as assessed by the grip strength test.
- E No significant differences in body weight between KO/HTT<sub>SD</sub> (n = 32), KO/HTT<sub>SA</sub> (n = 24), and KO mice (n = 24) at different postnatal timepoints.
- F Distance travelled during the open-field test by WT (n = 17), KO (n = 20), KO/HTT<sub>SD</sub> (n = 19), and KO/HTT<sub>SA</sub> mice (n = 17) at P35 and P55. Results are expressed in meters.
- G, H Results of 24-h Phenorack monitoring to measure the spontaneous activity of 12 WT, 12 KO, 8 KO/HTT<sub>SD</sub>, and 7 KO/HTT<sub>SA</sub> mice at P45 (G). Distance travelled at 8 AM versus 8 PM (H).

Data information: Kruskal–Wallis test with Dunn's comparison; \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, ns = not significant. Data are presented as the means  $\pm$  SEM.



Figure EV2.



### Figure EV3. Acute and chronic FK506 treatment of WT, HTT<sub>SA</sub>, and Mecp2 KO mice.

- A Western blot of HTT S421 phosphorylation. We treated WT and HTT<sub>SA</sub> 30-day-old mice intraperitoneally with FK506 (5 mg/kg) or vehicle and analyzed brain extracts for endogenous HTT phosphorylation by Western blotting, 2 h after administration (WT FK506 = 3, WT Vehicle = 6, HTT<sub>SA</sub> FK506 = 3, HTT<sub>SA</sub> Vehicle = 3). The relative protein level of phospho-HTT was normalized on total HTT protein level and is presented as the ratio. Data are presented as the means  $\pm$  SEM, \**P* < 0.05, Mann-Whitney test.
- B Western blot of HTT S421 phosphorylation. We treated WT, HTT<sub>SA</sub>, and *Mecp2* KO 30-day-old mice intraperitoneally with FK506 (5 mg/kg) or vehicle chronically for 20 days and analyzed brain extracts for endogenous HTT phosphorylation by Western blotting, 2 h after the last administration (KO FK506 n = 2, KO Vehicle n = 2, WT FK506 = 2, WT Vehicle = 2, HTT<sub>SA</sub> FK506 = 2, HTT<sub>SA</sub> Vehicle = 2).
- C Western blot of HTT and Mecp2. We treated WT and Mecp2 KO 30-day-old mice intraperitoneally with FK506 (5 mg/kg) or vehicle chronically for 20 days (WT FK506 = 3, WT Vehicle = 3, HTT<sub>SA</sub> FK506 = 3, HTT<sub>SA</sub> Vehicle = 3). Data are presented as the means  $\pm$  SEM, ns = not significant, Mann–Whitney test.

Source data are available online for this figure.

# Figure EV4. Absence of cytotoxicity and increased neuronal activity after chronic FK506 treatment.

- A FK506 did not increase the levels of cleaved caspase-3, as shown by immunostaining. Representative images of Striatum labeled with Casp3 (green) and DAPI of 50day-old Mecp2 KO mice treated with FK506 (right panel) or vehicle (left panel) for 20 days.
- B Absence of apoptosis in the dorsal striatum of *Mecp2*-knockout mice treated chronically with FK505 or with Vehicle. TUNEL assay was used to detect potential apoptotic neurons in 50-day-old *Mecp2*-deficient mice (*n* = 3 for each stage). Left panel: detection of cells containing fragmented DNA after desoxyribonuclease I treatment of a representative dorsal striatum section originating from one *Mecp2*-knockout and one wild-type animal. No labeled cells were visible after counting all striatum sections of 50-day-old *Mecp2*-knockout mice.
- C FK506 tends to increase the number of cFos-positive neurons. Representative images of striatum labeled with cFos (green) of 50-day-old KO mice treated with FK506 (right panel) or vehicle (middle panel) for 20 days.
- D Vertical scatter plot representing the number of cFos positive neurons counted in the dorsal striatum of P50 Wt placebo (n = 2)-, KO placebo (n = 3)-, and KO FK506 (n = 3)-treated mice in a picture of length 650  $\mu$ m and width 445  $\mu$ m. Each dot represents the mean of three successive sections.



Figure EV4.



## Figure EV5. Breathing patterns in 55-day-old WT, KO Vehicle, KO FK506, KO/HTT<sub>SD</sub>, and KO/HTT<sub>SA</sub> mice.

Left: Typical plethysmographic recordings of breathing (inspiration upward) performed in quiet, 55-day-old WT, KO Vehicle, KO FK506, KO/HTT<sub>SD</sub>, and KO/HTT<sub>SD</sub>