## 1 SUPPLEMENTARY FIGURES

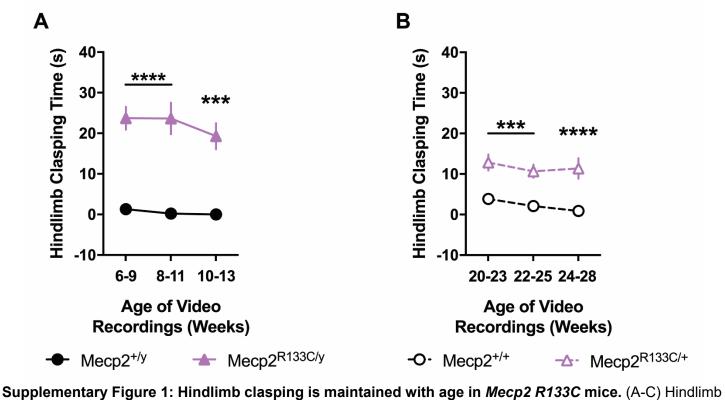
2 <u>**Title:**</u> Profiling beneficial and adverse effects of MeCP2 overexpression in a hypomorphic Rett syndrome mouse

3 model

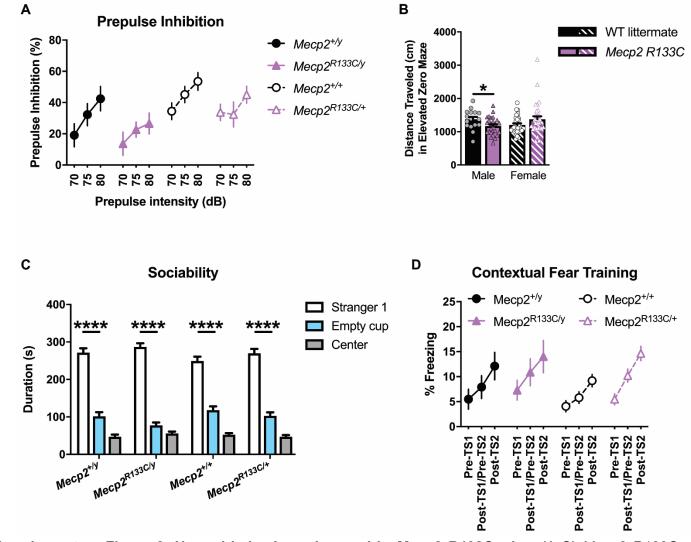
- 4 **<u>Running Title:</u>** MeCP2 overexpression in a hypomorphic Rett syndrome mouse
- 5 **Keywords:** Rett syndrome; MeCP2; *MECP2* Duplication syndrome; Neurodevelopment; Mouse model

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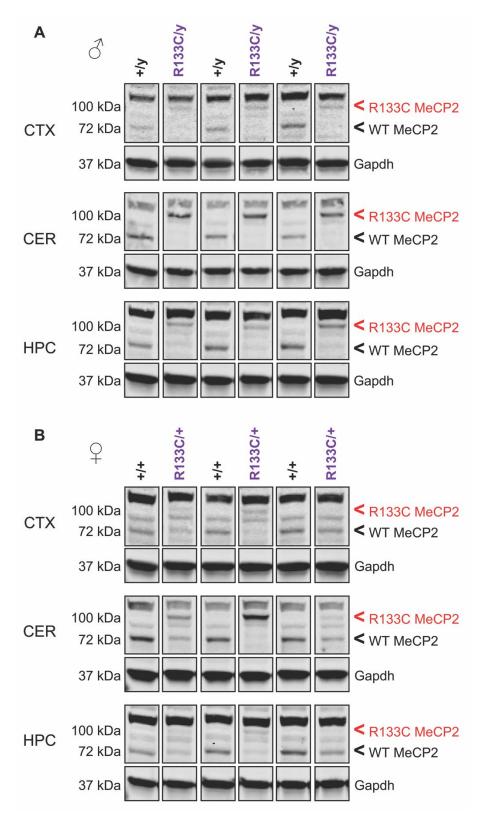
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Supplementary Figure 1: Hindlimb clasping is maintained with age in *Mecp2 R133C* mice. (A-C) Hindlimb clasping was measured across age for male  $Mecp2^{R133C/y}$  (6-13 weeks old) and female  $Mecp2^{R133C/+}$  (20-28 weeks old). n=16-30 per genotype in males, n=31-32 per genotype in females. 2-way ANOVA with Sidak's posthoc. \*within-genotype comparison. ns (not significant), \*\*\*p<0.001, \*\*\*\*p<0.0001. WT = closed or open black circles. *Mecp2 R133C* = closed or open purple triangles.

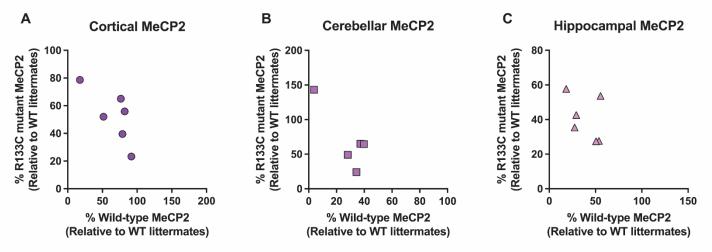


Supplementary Figure 2: Normal behaviors observed in Mecp2 R133C mice. (A-C) Mecp2 R133C mice 31 32 displayed normal sensorimotor gating in pre-pulse inhibition (PPI) across several pre-pulse intensities (A), distance traveled in the elevated zero maze (B) and social behavior (C) assays, and spent more time with 33 Stranger 1 mouse than the empty cup in the 3-chamber task. (D) On training day of contextual fear conditioning, 34 male *Mecp2*<sup>*R*133C/y</sup> and female *Mecp2*<sup>*R*133C/+</sup> responded normally to the aversive stimuli. n=13-29 per genotype in 35 36 males, n=18-32 per genotype in females. 2-way ANOVA with t-test post-hoc, or unpaired t-test. \*within-genotype comparison. \*\*\*\*p<0.0001. WT = filled or patterned black bars or closed or open black circles. Mecp2 R133C = 37 filled or patterned purple bars or closed or open purple triangles. Male = filled, closed. Female = patterned, open. 38



Supplementary Figure 3: WT and *R133C* mutant MeCP2 proteins are detected in brain samples via Western Blotting. (A-B) Representative images of Western blots and distinct bands are indicated for the *R133C* mutant (red arrow) and WT MeCP2 (black arrow), as well as the Gapdh control, in male (3) and female (9)

- 42 Mecp2 R133C mice and WT littermates across several brain regions. +/y or +/+ (WT), R133C/y or R133C/+
- 43 (*Mecp2 R133C*). CTX (cortex), CER (cerebellum), HPC (hippocampus).

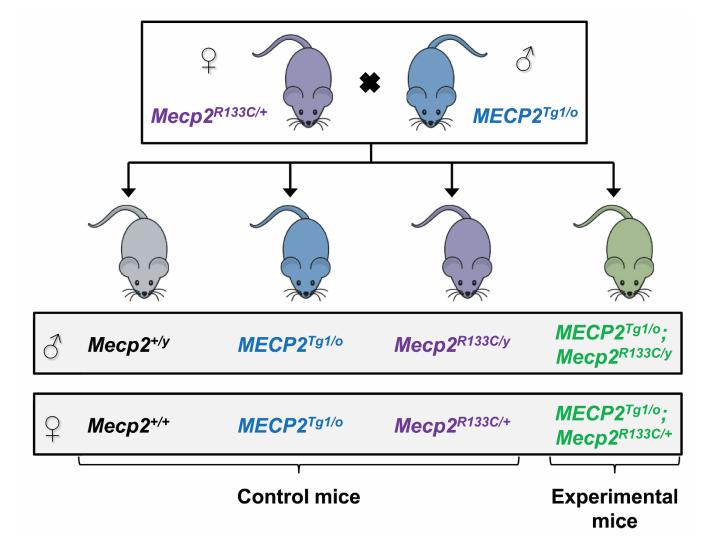


44 Supplementary Figure 4: Variability of WT and *R133C* mutant MeCP2 expression in *Mecp2*<sup>*R133C/+*</sup> mice.

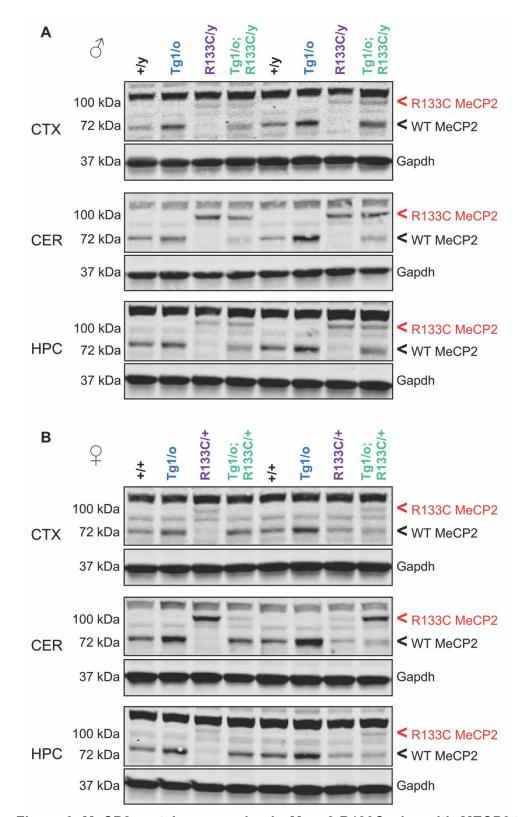
The relationship of WT MeCP2 and the R133C mutant MeCP2 expression in the cortex, cerebellum and

46 hippocampus of *Mecp2*<sup>*R*133C/+</sup> mice is illustrated. Each point corresponds to a mouse subject, and the expression

- 47 of both forms of MeCP2 is normalized to the Gapdh control and presented as percentage expression relative to
- 48 MeCP2 expression in WT littermates.



Supplementary Figure 5: Breeding scheme to generate *Mecp2 R133C* mice expressing an *MECP2* transgene. Male  $MECP2^{Tg1/o}$  mice (blue) were bred with female  $Mecp2^{R133C/+}$  (purple) animals to introduce the human *MECP2* transgene and generate the experimental mice  $MECP2^{Tg1/o}$ ;  $Mecp2^{R133C/y}$  and  $MECP2^{Tg1/o}$ ;  $Mecp2^{R133C/+}$  (green). Wild-type (gray),  $MECP2^{Tg1/o}$ , and  $Mecp2^{R133C/y}$  or  $Mecp2^{R133C/+}$  littermates were used as controls.

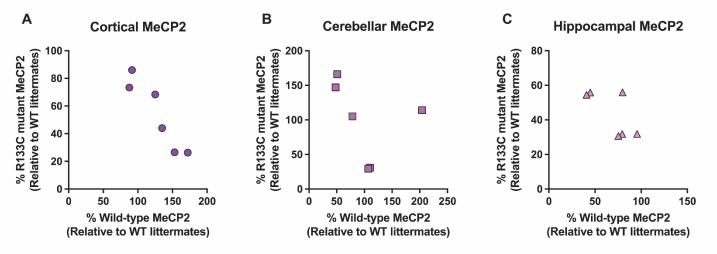


54 Supplementary Figure 6: MeCP2 protein expression in *Mecp2 R133C* mice with *MECP2* transgene. (A-B)

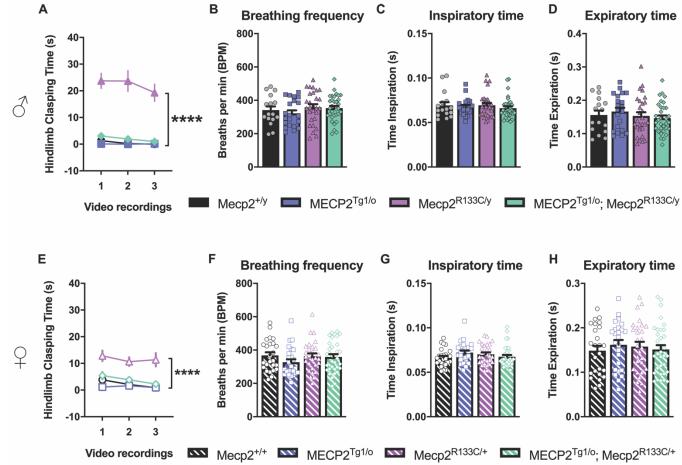
55 Representative images of Western blots showing distinct bands for the *R133C* mutant (red arrow) and WT

56 MeCP2 (black arrow), as well as the Gapdh control, in male ( $\bigcirc$ ) and female ( $\bigcirc$ ) mice across several brain regions.

- 57 +/y or +/+ (WT), Tg1/o (*MECP2<sup>Tg1/o</sup>*), R133C/y or R133C/+ (*Mecp2 R133C*), Tg1; R133C/y or Tg1; R133C/+
- 58 (*Mecp2 R133C* with *MECP2* transgene). CTX (cortex), CER (cerebellum), HPC (hippocampus).



Supplementary Figure 7: Variability of WT and *R133C* mutant MeCP2 expression in *MECP2<sup>Tg1/o</sup>*; *Mecp2<sup>R133C/+</sup>* mice. The relationship of WT MeCP2 and the *R133C* mutant MeCP2 expression in the cortex, cerebellum and hippocampus of  $Mecp2^{R133C/+}$  mice with the *MECP2* transgene is illustrated. Each point corresponds to a mouse subject, and the expression of both forms of MeCP2 is normalized to the Gapdh control and presented as percentage expression relative to MeCP2 expression in WT littermates.



64 Supplementary Figure 8: Hindlimb clasping and respiratory function parameters in Mecp2 R133C mice with MECP2 transgene. (A, E) Hindlimb clasping is reversed across all ages in MECP2<sup>Tg1/o</sup>; Mecp2<sup>R133C/y</sup> and 65 *MECP2<sup>Tg1/o</sup>; Mecp2<sup>R133C/+</sup>* mice. Video recording numbers represent different ages: 6-9, 8-11 or 10-13 weeks old 66 67 in males; 20-23, 22-25 or 24-28 weeks old in females. (B-D, F-H) No differences in other respiratory parameters between genotypes in males or females, including breathing frequency, and times of inspiration or expiration, 68 were observed. n=16-32 per genotype in males, n=25-40 per genotype in females. 1- or 2-way ANOVA with 69 Tukey's post-hoc tests. \*\*\*\*p<0.0001. WT = black bars or circles. *MECP2<sup>Tg1/o</sup>* = blue bars or squares. *Mecp2* 70 R133C = purple bars or triangles. *MECP2<sup>Tg1/o</sup>*; *Mecp2* R133C = green bars or diamonds. Male (3) = filled bars 71 72 or closed circles. Female ( $\mathcal{Q}$ ) = patterned bars or open circles.