

Figure S1 Absence of histopathology in P0 N-CKO tissue. **(A)** Nissl stain of P0 N-CKO brain. Scale bar= 1 mm. **(B)** GFAP immunostaining. Scale bar= 1 mm. **(C)** GFAP and AC3 immunoreactivity in P10 spinal cords. N-CKO spinal cord exhibits gliosis and apoptosis. Arrows highlight activated caspase-3-positive nuclei in ventral horn (VH). DH: dorsal horn; VH: ventral horn. Scale bar= 100 μ m.

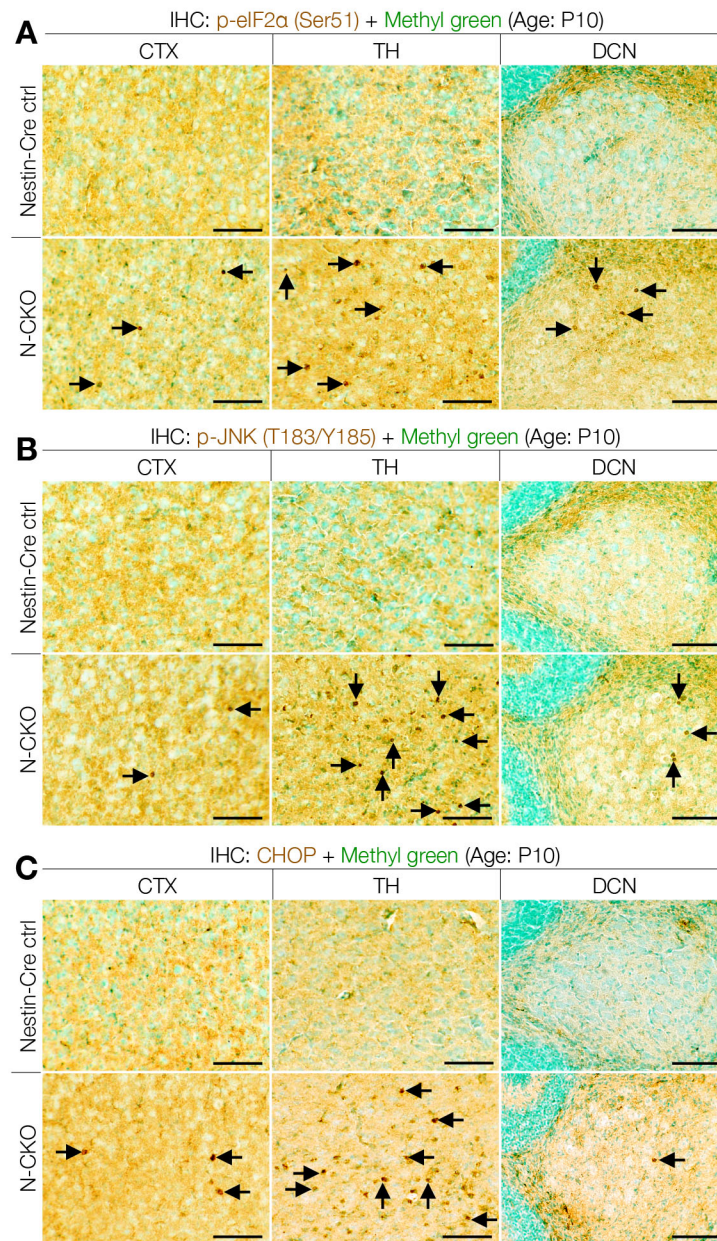


Figure S2 N-CKO CNS tissue exhibits markers of ER-stress in susceptible sensorimotor regions. Immunostaining for **(A)** phospho-eIF2 α (Ser51), **(B)** phospho-JNK (T183/Y185), and **(C)** nuclear-translocated CHOP is increased in vulnerable regions of N-CKO brains. Arrows highlight positive cells. Scale bar= 100 μ m.

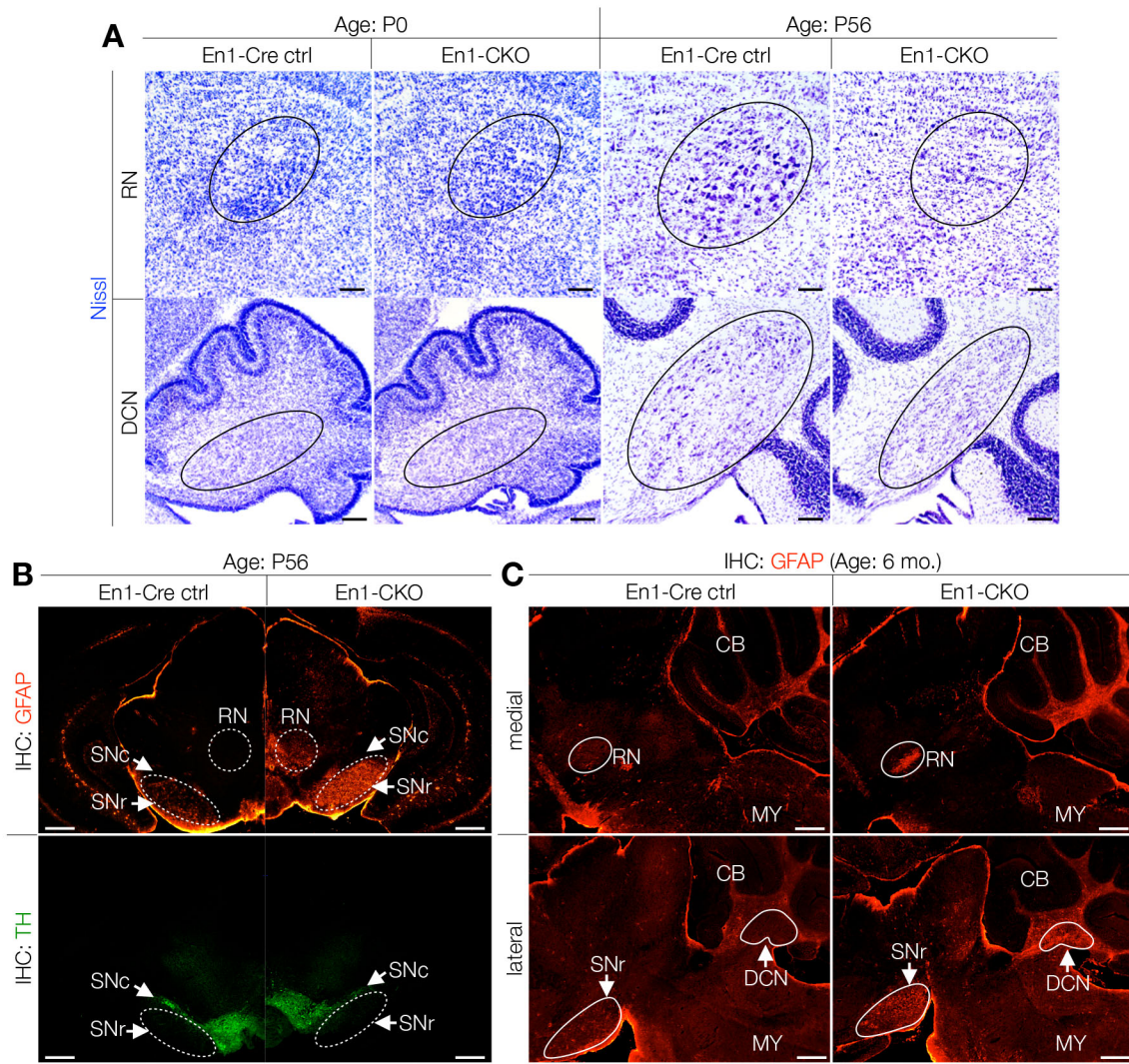


Figure S3 Neurodegeneration in En1-CKO mutants. **(A)** Nissl staining of En1-CKO and control brain sections. En1-CKO brains appear normal at P0 but show profound neuronal loss in RN and DCN at P56. Scale bar= 100 μ m. **(B)** GFAP and tyrosine hydroxylase (TH) immunostaining (P56). Gliosis is present selectively in RN and SNr of En1-CKO. SNc: substantia nigra pars compacta. Scale bar = 500 μ m. **(D)** GFAP Immunoreactivity in 6-month En1-CKO and control brain sections. Gliosis remains confined tp RN, SNr, and DCN, with no spread to other Cre-expressing structures. Scale bar= 100 μ m.

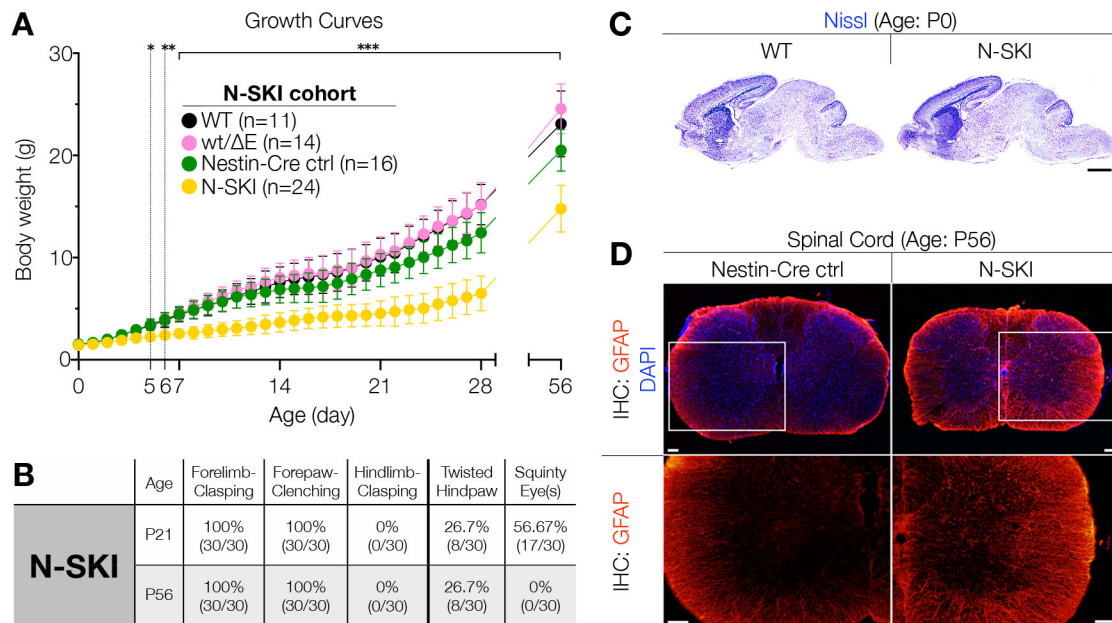


Figure S4 Characterization of N-SKI mice. **(A)** Growth curves. N-SKI mice become significantly lighter from P5 onward. 2-Way Anova, Mean (SD), * $P < 0.05$; ** $P < 0.01$; *** $P < 0.001$. **(B)** The frequency of N-SKI mutants showing various behavioral abnormalities at P21 and P56. **(C)** Nissl staining of N-SKI and control brains (P0). Scale bar= 1 mm. **(D)** GFAP immunoreactivity in N-SKI and control P56 spinal cords. Mild gliosis is present N-SKI tissue. Scale bar= 100 μm .

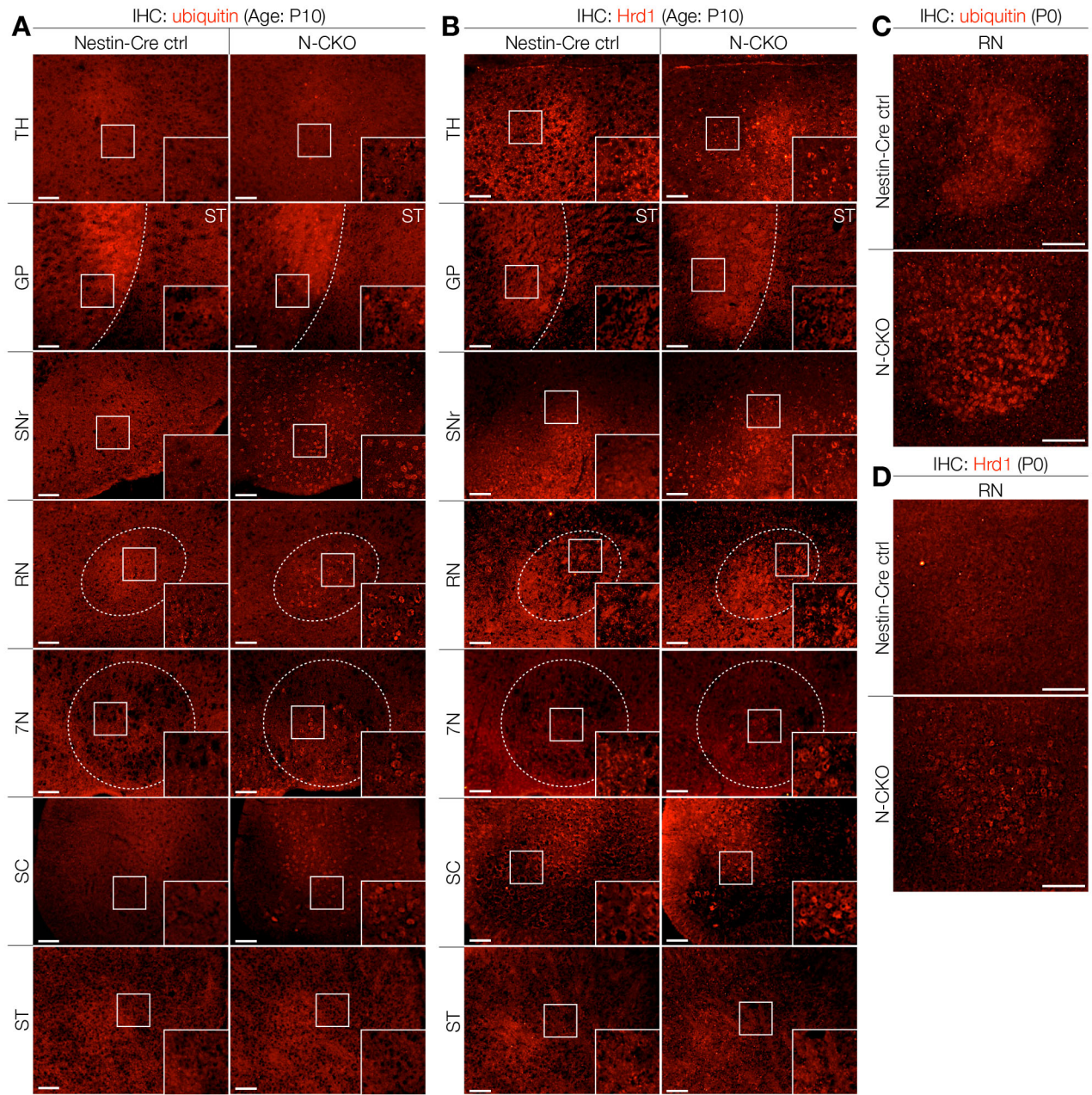


Figure S5 Region selective abnormalities of ubiquitin and Hrd1 in N-CKO mice. For both molecules, the lack of abnormality in striatum illustrates that these molecular abnormalities are isolated to the structures exhibiting gliosis and neurodegeneration (**A**) Perinuclear accumulation of ubiquitin immunoreactivity (P10). Scale bar= 100 μ m. (**B**) Perinuclear accumulation of Hrd1 immunoreactivity (P10). Scale bar= 100 μ m. Abnormal perinuclear accumulation of ubiquitin (**C**) and Hrd1 (**D**) can be detected in RN of N-CKO as early as at P0. Scale bar= 100 μ m.

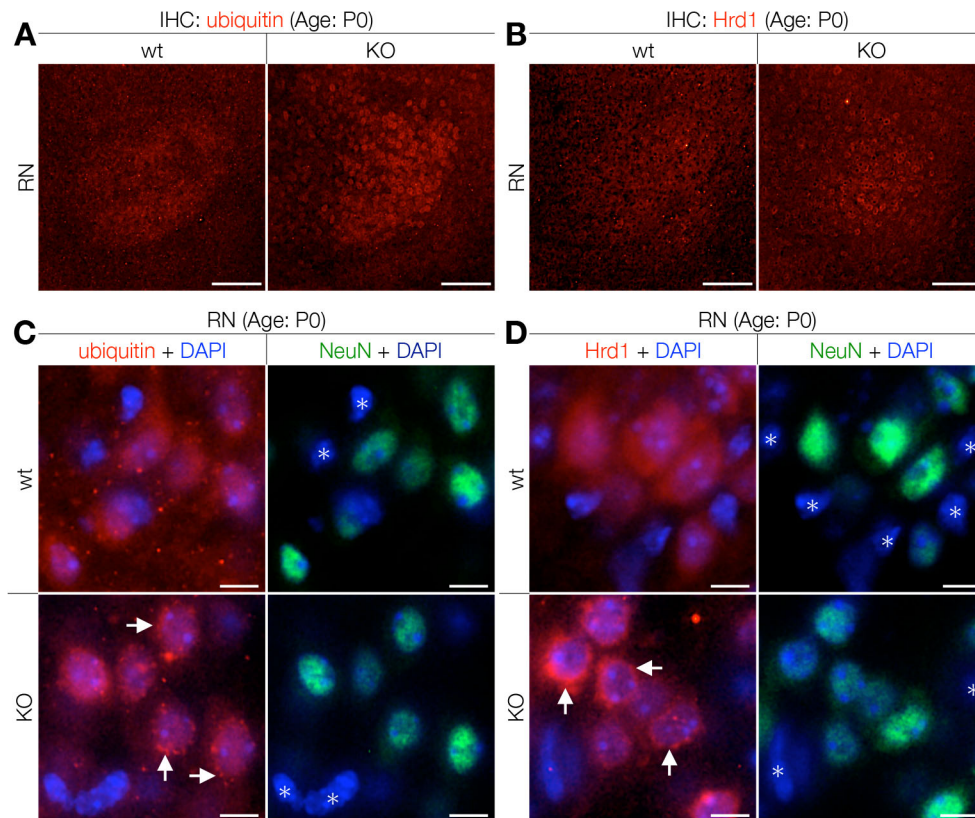


Figure S6 Abnormal perinuclear accumulation of ubiquitin (**A**) and mislocalization of Hrd1 (**B**) in the RN of torsinA constitutive knockout (KO) at P0. Scale bar= 100 μ m. (**C**) Only NeuN-positive cells in RN of torsinA-null mice (KO) show perinuclear ubiquitin-accumulation. Asterisks mark non-neuronal cells. Scale bar= 100 μ m. (**D**) Hrd1 expression is neuron-specific and only NeuN-positive cells in RN of KO show abnormal Hrd1 mislocalization. Asterisks mark non-neuronal cells. Scale bar= 10 μ m.

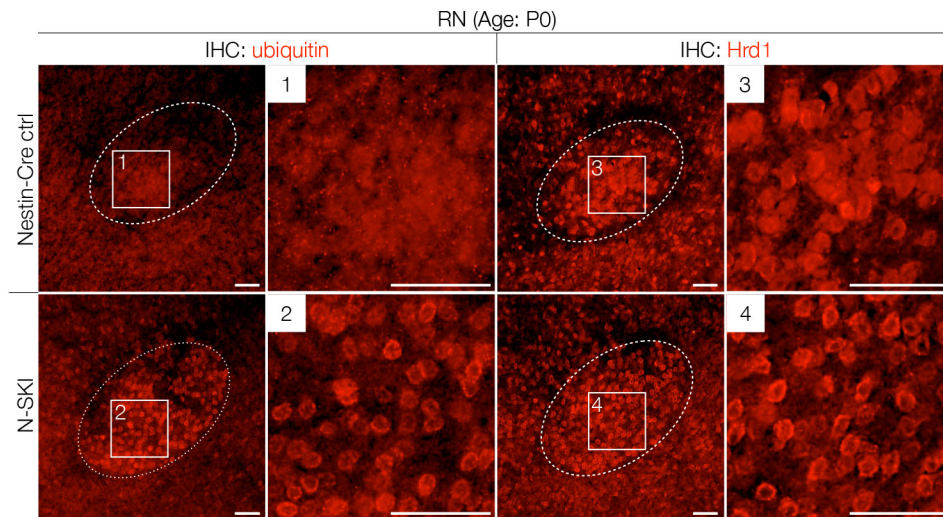


Figure S7 Region selective abnormalities of ubiquitin and Hrd1 in N-SKI mice. Immunohistochemistry of ubiquitin and Hrd1 in P0 RN. Abnormal perinuclear ubiquitin accumulation and Hrd1 mislocalization show in RN of N-SKI at P0. Scale bar= 100 μ m.