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2 **Title:** Optimization of systemic AAV9 gene therapy in Niemann-Pick disease type C1 mice

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4 **Running Title:** Systemic AAV9 gene therapy in *Npc1<sup>m1N</sup>* mice

5  
6 **Summary Blurb:** Systemic AAV9-*hNPC1* gene therapy in null *Npc1<sup>m1N</sup>* mice at higher doses or with  
7 earlier administration and treatment of hypomorphic *Npc1<sup>I1061T</sup>* mice delays disease progression and  
8 increases lifespan.

9  
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44 **Abstract (175 words):**

45  
46 Niemann-Pick disease, type C1 (NPC1) is a rare, fatal neurodegenerative disorder caused by pathological  
47 variants in *NPC1*, which encodes a lysosomal cholesterol transport protein. There are no FDA approved  
48 treatments for this disorder. Both systemic and central nervous system delivery of AAV9-*hNPC1* have  
49 shown significant disease amelioration in NPC1 murine models. To assess the impact of dose and  
50 window of therapeutic efficacy in *Npc1<sup>mlN</sup>* mice, we systemically administered three different doses of  
51 AAV9-*hNPC1* at 4 weeks old and the medium dose at pre-, early, and post-symptomatic timepoints.  
52 Higher vector doses and treatment earlier in life were associated with enhanced transduction in the  
53 nervous system and resulted in significantly increased lifespan. Similar beneficial effects were noted after  
54 gene therapy in *Npc1<sup>11061T</sup>* mice, a model that recapitulates a common human hypomorphic variant. Our  
55 findings help define dose ranges, treatment ages, and efficacy in severe and hypomorphic models of  
56 NPC1 deficiency and suggest that earlier delivery of AAV9-*hNPC1* in a pre-symptomatic disease state is  
57 likely to yield optimal outcomes in individuals with NPC1.

58  
59 **Introduction:**

60  
61 Niemann-Pick disease, type C (NPC) is a rare, fatal neurodegenerative disease with an incidence of ~1 in  
62 100,000 live births [1]. This autosomal recessive lysosomal storage disorder is marked by unesterified  
63 cholesterol and sphingolipid accumulation in the lysosome, the latter of which is especially prominent in  
64 neural tissue. At least 95% of NPC1 individuals have disease associated variants in the integral  
65 membrane-bound NPC1 protein located in the lysosome (NPC1 disease, OMIM #257220). The remaining  
66 individuals have pathological variants in the NPC2 protein (NPC2 disease, OMIM #607625), a soluble  
67 lysosomal protein that transfers unesterified cholesterol to NPC1 [2]. Clinical presentation of both forms  
68 of the disease are similar, which is in accordance with previous research demonstrating that NPC1 and  
69 NPC2 work together to facilitate egress of cholesterol and likely other lipids from the lysosome [1, 3-6].  
70 Disease severity and onset is highly variable, affecting infants, children, and adults; however, the classical  
71 presentation of NPC1 is most often observed in school-age children and typically includes progressive  
72 cerebellar ataxia, vertical supranuclear gaze palsy, gelastic cataplexy, motor deficits, cognitive  
73 impairment as well as visceral manifestations like hepatosplenomegaly [7, 8]. Miglustat, a  
74 glycosphingolipid synthesis inhibitor, is approved for treatment for NPC outside the US and though other  
75 potential therapeutics have advanced through trial or to expanded access protocols, none have secured  
76 regulatory approval [9-12]. While small molecule therapeutics provide some amelioration, all fail to  
77 address the root cause of the disorder – the absence of the NPC1 protein that causes subsequent morbidity  
78 and mortality. Gene therapy can provide the replacement of the dysfunctional NPC1 protein to treat the  
79 disease [13, 14]. Given the irreversible nature of neurodegeneration and lethality, NPC individuals remain  
80 in dire need of effective treatments that a durable gene therapy might provide.

81  
82 Recent studies have demonstrated that gene therapy, especially those using adeno-associated viral (AAV)  
83 vectors, can provide treatments for monogenic and rare diseases. AAVs are already approved for  
84 treatment of Spinal Muscular Atrophy (SMA1; onasemnogene abeparvovec-xioi [15]) and RPE65  
85 mutation associated retinal dystrophy (voretigene neparvovec-rzyl [16]) [17-20]. AAV serotypes are well-  
86 characterized and increasingly used in clinical trials for monogenic diseases [21-24]. Vectors that use  
87 AAV serotype 9, or AAV9, are well documented to cross the blood-brain barrier (BBB) and transduce  
88 cells of the central nervous system (CNS) [25-27]. Given the devastating neurological impact of NPC1, a  
89 gene therapy targeting the CNS is imperative. Of note, AAV9 also transduces multiple other organ  
90 systems, including the liver and peripheral nerves, that are implicated in NPC1 disease [7, 8, 28-30].

91  
92 We and others have previously demonstrated AAV9 vectors can effectively improve survival and delay  
93 NPC1 disease progression in a severe NPC1 murine model (*Npc1<sup>mlN</sup>*) [21, 31-35]. We will use single

94 allele notation to indicate homozygosity. Both direct CNS administration (intracerebroventricular or  
95 intracisternal magna) [32, 34] and systemic administration (retro-orbital or intracardiac) [21, 31] have  
96 successfully ameliorated disease in these mice. Greater success has been noted when using dual routes of  
97 CNS administration or administering higher doses of vector to the CNS [34, 35]. Independent studies  
98 have demonstrated that ubiquitous promoters provide greater disease correction in *Npc1<sup>m1N</sup>* mice as  
99 compared to neuron-specific promoters [21, 35] and further optimization studies have highlighted the  
100 therapeutic potential of novel capsids to improve CNS transduction [33].

101  
102 While there is a robust foundation for gene therapy studies largely focused on neonatal intervention in  
103 severely affected *Npc1<sup>m1N</sup>* mice, further studies using variable times of delivery and mouse models that  
104 recapitulate attenuated forms of the disease have not been fully explored. Nearly all preclinical studies in  
105 *Npc1<sup>m1N</sup>* murine models have targeted neonates [31, 32, 34, 35]. In our previous work [21, 33], AAV9  
106 vectors were administered at pre-symptomatic, weaning age (4 weeks old), but the question has remained  
107 of whether late(r) intervention can still be effective following early or post-symptomatic diagnoses in  
108 individuals. Previous clinical studies for another rare pediatric genetic disorder, aromatic L-amino acid  
109 decarboxylase deficiency, suggest that AAV gene therapy is universally beneficial, but treatment at a  
110 younger age was associated with greatest therapeutic effects (NCT01395641, NCT02926066) [36].  
111 Intervention prior to clinical onset of neurologic symptoms in NPC is currently challenging because  
112 newborn screening for NPC1 is not yet included on the Recommended Uniform Screening Panel (RUSP)  
113 and the average diagnostic delay remains ~4.1 years [37, 38]. Although NPC1 is on the American College  
114 of Obstetricians and Gynecologists (ACOG) and American College of Medical Genetics and Genomics  
115 (ACMG) suggested carrier screening panel to identify couples at risk for affected pregnancies, it is not  
116 routinely screened for. Early intervention prior to neurologic onset might be possible in familial cases and  
117 after diagnosis when there is infantile presentation with fetal ascites and liver disease [39-41]. More than  
118 600 pathogenic or likely pathogenic NPC1 variants have been described, most of which are missense  
119 mutations [42-44]. One of the most prevalent variants results in a missense mutation in the NPC1  
120 p.I1061T protein, causing NPC1 to misfold and undergo endoplasmic reticulum associated degradation  
121 (ERAD) [45, 46]. A knock-in, hypomorphic *Npc1<sup>I1061T</sup>* allele was generated to recapitulate the human  
122 disorder [47, 48]: *Npc1<sup>I1061T</sup>* mice have a slightly protracted disease course compared to the *Npc1<sup>m1N</sup>*  
123 mouse model that has a premature stop codon in the *Npc1* gene resulting in production of truncated, non-  
124 functional NPC1 protein [47]. Using *Npc1<sup>I1061T</sup>* mice, we investigated and confirmed that residual NPC1  
125 protein with compromised stability did not interfere with the efficacy of gene therapy.

126  
127 Here we build on our previous work, using systemic administration of an AAV9-elongation factor 1 $\alpha$   
128 (shortened)-*hNPC1* (AAV9-EF1a(s)-*hNPC1*) vector to treat various mouse models of NPC1 at timepoints  
129 later in disease progression. We further examine the therapeutic efficacy of this vector across different  
130 doses and in *Npc1<sup>I1061T</sup>* mice. In aggregate, our results provide foundational preclinical data for the  
131 advancement of AAV9-EF1a(s)-*hNPC1* as a disease modulating therapy for individuals with NPC1  
132 deficiency.

133  
134  
135 **Results:**

136  
137 **Mice treated with higher doses of AAV9-EF1a(s)-*hNPC1* gene therapy showed increased survival**  
138 **and delayed disease progression.**

139 All mice received a retro-orbital injection of AAV9-EF1a(s)-*hNPC1*, herein referred to as AAV9, and  
140 every figure panel following contains the data of 4-28 mice per group (Table S1). To compare the  
141 efficacy of AAV9 at varying doses, mice were injected at 4 weeks old (weaning, postnatal day 26-28)  
142 with a low, medium, or high dose ( $7.87 \times 10^{12}$  vector genomes/kg,  $1.28 \times 10^{14}$  vg/kg, or  $3.06 \times 10^{14}$  vg/kg,  
143 respectively) Based on a Log-Rank, Mantel Cox test of survival with application of Bonferroni's  
144 correction, all treated cohorts showed improvement over saline injected mice. Mice treated with high and

145 medium doses survived longer than low dose and saline-only injected mice (34.6, 21.5, 11.4, and 10.6  
146 weeks median survival, respectively,  $P < 0.0001$ ) (Fig 1A, B). Of note, the low dose injected mice also had  
147 improved survival compared to the saline-only group (11.4 vs 10.6 weeks,  $P = 0.0049$ ).

148  
149 *Npc1<sup>mIN</sup>* mice exhibit marked weight loss starting at about 6 weeks old; therefore, the week that mice  
150 reached their peak weight and the change in weight from 6 to 9 weeks of age was analyzed. Mice  
151 receiving the low dose reached peak weight at  $6.9 \pm 0.6$  weeks, similarly to saline injected controls ( $6.8 \pm$   
152  $0.7$  weeks). Both groups reached peak weight earlier than either medium dose mice ( $11.3 \pm 3.9$  weeks) or  
153 high dose mice ( $14.8 \pm 3.1$  weeks) (Fig 1C, Kruskal-Wallis Test with Dunn's multiple comparisons test).  
154 Longitudinal weight data (Fig S1A, B) demonstrates that mice maintain weight and survive longer as the  
155 dose of AAV9 increases. Both saline and low dose mice lost weight similarly between 6 and 9 weeks  
156 ( $-14.6\% \pm 6.8\%$  and  $-9.0\% \pm 11.7\%$ , respectively) (Fig 1D). Notably, medium ( $11.2\% \pm 13.1\%$ ) and high  
157 dose ( $15.3\% \pm 10.9\%$ ) treated mice showed significant differences from both saline and low dose, gaining  
158 weight at a similar trajectory to *Npc1<sup>+/+</sup>* mice ( $5.6\% \pm 4.1\%$ ) between 6 and 9 weeks of age (One-way  
159 ANOVA with Tukey's multiple comparisons test).

160  
161 To determine the effect of dose on phenotypic progression, five parameters indicative of neurological  
162 phenotype (hindlimb clasp, motor function, kyphosis, grooming, and a balance-ledge test) were assessed  
163 in the mice at 3-week intervals from weeks 6-18 [49]. Phenotype assessment shows most significant delay  
164 of disease onset and progression in high dose mice, but there was still notable delay in medium dose  
165 mice. Two-way ANOVA tests with mixed effects analysis were used to estimate how the mean phenotype  
166 score changes according to the time of assessment and treatment group. Low dose mice follow the vehicle  
167 trajectory of progression (Fig 1E). Between 6 and 9 weeks, all treatment groups were significantly  
168 delayed in phenotype onset from vehicle treated, while high dose treated mice were like *Npc1<sup>+/+</sup>*.  
169 Between weeks 9 and 12, mice treated with high and medium doses had significantly lower composite  
170 scores than low dose and saline injected mice ( $P < 0.0001$ , two-way ANOVA mixed-effects analysis with  
171 Tukey's multiple comparisons test, results presented in Fig S2A). Comparing weeks 15 and 18, high dose  
172 treated mice had significantly lower composite scores than medium dose treated ( $P < 0.01$ , two-way  
173 ANOVA mixed-effects analysis with Tukey's multiple comparisons test), and high dose treated mice  
174 were not significantly different from *Npc1<sup>+/+</sup>* but still trended upwards.

175  
176  
177 **Higher doses of AAV9 result in greater viral transduction in *Npc1<sup>mIN</sup>* mice.**

178  
179 To evaluate the efficacy of vector transduction in various tissues, droplet digital PCR (ddPCR) was  
180 performed using cerebrum (Fig 2Ai, ii, 2C) and liver (Fig 2Bi, ii) to determine *hNPC1* copy number at 10  
181 weeks of age or the humane endpoint/survival. At 10 weeks old, the typical age for humane endpoint  
182 without therapeutic intervention in our colony, there were higher copy numbers in the high dose cerebrum  
183 compared to all other treatment groups (Fig 2Ai). There was only a significant predictive relationship  
184 between *hNPC1* copy number and lifespan in the cerebrum within the medium dose group (Fig 2Aii), but  
185 there was a strong correlation between copy number in cerebrum and increased survival when all treated  
186 mice are grouped compared to saline-injected mutants (Fig 2C). Of note, two mice had exceptionally and  
187 unexpectedly high CNV in the cerebrum in the medium dose treated group, and seem to drive a  
188 significant relationship between CNV and lifespan within the medium dose group, but not when all  
189 treated mice are grouped together (Fig 2Aii, C). In the liver, there were higher copy numbers in the high  
190 dose mice compared to other treatment groups at 10 weeks (Fig 2Bi), but there was no predictive  
191 relationship between copy number and increased survival within each treatment group (Fig 2Bii).  
192 Additional analysis of 10-week-old mice was performed to understand copy number variation across  
193 different organs (spleen, kidney, lung, muscle, cerebellum, brain stem). Overall, an increased dose  
194 (particularly high dose) of gene therapy led to greater copy numbers (Fig 2D).

195

196 Western blots were performed to determine the level of NPC1 protein present in brain and liver tissue in  
197 10-week old *Npc1<sup>miN</sup>* mice where the only protein present would be a result of vector transduction (Fig  
198 2C, F, & representative blot Fig S5). NPC1 levels assessed in the cerebrum did not reveal significant  
199 levels of protein for any of the doses, though cerebrum from high dose treated mice revealed very low,  
200 albeit detectable levels of protein (Fig 2C). In livers, NPC1 protein was present in medium and high dose  
201 treated mice. High dose treated mice displayed a greater level of NPC1 protein compared to *Npc1<sup>+/+</sup>* (not  
202 considered significant). Low dose treated mice did not have appreciable levels of detectable protein (Fig  
203 2F).

204  
205

### 206 **Higher doses reduce pathology in the brain and liver and preserve cerebellar Purkinje neurons of** 207 **10-week-old mice.**

208

209 Immunofluorescent staining of the brain and immunohistochemical staining of the liver was performed to  
210 assess cholesterol storage in the cerebellum, health of the cerebellum, and inflammation in age-matched  
211 10-week-old mice. Parallel assessments of GFAP (astrocytes), calbindin D (Purkinje neurons), and CD68  
212 (microglia) and NPC1 proteins via western blot were used for quantification. Unesterified cholesterol  
213 storage, as ascertained by filipin labeling, appears modestly reduced with high dose gene therapy when  
214 compared to lower doses or saline in 10-week-old *Npc1<sup>miN</sup>* mice (Fig 3A). Neuroinflammation as  
215 indicated by GFAP labeling of astrocytes and western blot of cerebellum is reduced (not statistically  
216 significant) with increasing dose and appears to correlate with higher copy numbers of *hNPC1* (Fig 3A,  
217 C, & representative blot Fig S6). Purkinje neuron survival appears to increase in the cerebellum as gene  
218 therapy dose increases but protein levels in treated mice are significantly less than in *Npc1<sup>+/+</sup>* mice;  
219 Purkinje neuron preservation seems to be correlated with copy number of *hNPC1* (Fig 3A, D,  
220 representative blot Fig S7). Anterior to posterior loss of Purkinje neurons is concurrent with previous  
221 findings in NPC1 transcriptomics of the cerebellum (Fig 3A; [50]). Higher CNV also correlates with  
222 reduced neuroinflammation, as evidenced by decreased immunostaining of microglia in posterior lobules  
223 of the cerebellum and apparent reduced CD68 protein levels in the cerebellum of high dose mice (Fig 3B,  
224 E respectively, representative blot Fig S6).

225

226 In the cerebellum, ddPCR of the *hNPC1* gene (Fig 3F) reveals an apparent linear relationship between  
227 increasing copy number and survival in medium dose mice, though it is not significant in any other  
228 treatment group. There are two outliers in the medium dose mice (same two mice as seen in the cerebrum  
229 analysis) that when removed, mean the relationship between CNV and survival in this group is not  
230 significant. NPC1 levels were assessed in the cerebellum via western blot, demonstrating higher apparent  
231 levels of NPC1 as dose increased (Fig 3G, representative blot Fig S7).

232

233 NPC1 liver pathology includes an increased presence of myeloid cells (CD68+) due to increased lipid  
234 storage burden within these cells, many of which are macrophages like Kupffer cells (KC) (Fig 4A, B).  
235 Lipid laden myeloid cells are highly abundant in *Npc1<sup>miN</sup>* mice administered saline or low dose gene  
236 therapy. There is a dose-dependent effect on myeloid cell labeling, with percent CD68+  
237 immunohistochemical labeling decreasing with more effective, higher doses (Fig 4B, C). Fluorescent  
238 imaging also reveals decreases in cholesterol storage at high doses, demonstrated by diminished presence  
239 of myeloid cells (CD68) and a decrease in amount of filipin labeling (Fig 4A; specifically, inset for high  
240 dose liver with arrows indicating groups of cells without cholesterol storage). Overall, increased doses  
241 appear to reduce liver pathology.

242

243 Upon evaluation of the survival, weight curves, and phenotype data from the three main studies (dose, age  
244 at treatment and hypomorphic model), we observed the greatest disease amelioration with a high dose;  
245 thus, we carried out more detailed analysis and quantification of pathology in the dose cohorts. Age of

246 treatment and efficacy in the *Npc1*<sup>I1061T</sup> model data followed similar trends with respect to correlation  
247 between survival time and impact on pathology.

248

#### 249 **Alterations in sphingolipid distribution following gene therapy administration.**

250 In addition to cholesterol accumulation, multiple other lipid classes exhibit altered levels resulting from  
251 impaired NPC1 protein function [51, 52]. Mass spectrometry imaging was performed to evaluate the  
252 effect of gene therapy on lipid distribution following administration. A stark contrast in distribution was  
253 observed when comparing mutant and *Npc1*<sup>+/+</sup> mice, especially within the sphingolipid class (Figure S8).  
254 Gangliosides such as GM2 are known to accumulate within the brain in *Npc1*<sup>m1N</sup> mice [53]. *Npc1*<sup>+/+</sup> mice  
255 demonstrated little to no ganglioside accumulation within the brain, compared to the mutant mice which  
256 displayed high ganglioside accumulation, primarily in the frontal cortex and lobule X in the cerebellum.  
257 With increasing doses of gene therapy, a corresponding qualitative reduction in ganglioside accumulation  
258 was observed in both the cortex and cerebellum (Figure S8A). In contrast, the sphingolipid  
259 hexosylceramide (HexCer 46:4;O3) is increased in the cerebellum of *Npc1*<sup>+/+</sup> mice compared to the  
260 mutant. While the frontal cortex remains largely unchanged between doses, a change in lipid abundance is  
261 observed at differing AAV9 doses (Figure S8B). While hexosylceramide signal increased throughout the  
262 cerebellum, the rostral lobes of the cerebellum (lobules I-V) display a higher abundance compared to the  
263 rest of the cerebellum. Another sphingolipid, dihydroceramide (Cer 32:2;O3), was found to be increased  
264 in mutant mice compared to *Npc1*<sup>+/+</sup> mice. However, reduction in dihydroceramide accumulation is less  
265 apparent, though high and medium dose gene therapy do appear to impact this lipid (Figure S8C).

266

267

#### 268 **Treating mice at 4 weeks of age with AAV9-EF1a(s)-hNPC1 gene therapy significantly improves** 269 **survival and reduces disease progression compared to treatment at 6 or 8 weeks of age.**

270

271 *Npc1*<sup>m1N</sup> mice were treated with 1.28x10<sup>14</sup> gene copies/kg of AAV9-EF1a(s)-hNPC1 at 4 weeks (pre-  
272 symptomatic), 6 weeks (early-symptomatic), or 8 weeks (late-symptomatic) to determine the therapeutic  
273 window of efficacy for gene therapy. Comparison of survival between these groups (Fig 5A) shows that  
274 mice injected at 4 weeks had a significantly longer median survival (21.5 weeks) than saline injected mice  
275 (10.6 weeks) or mice treated at either 8 weeks (11.9 weeks) or 6 weeks (13.2 weeks). Bonferroni's  
276 correction factor was applied for 6 comparisons that lowered level of significance to P=0.0083 for this  
277 Log-rank Mantel Cox survival test (Fig 5B).

278

279 Week of peak weight and percent weight change between 6 and 9 weeks was also assessed to determine  
280 how gene therapy affected weight. Mice injected at 4 weeks old had a significantly later week of peak  
281 weight (11.3 ± 3.9 weeks) than saline injected mice or mice treated at 6 or 8 weeks of age (P <0.0001)  
282 (Fig 5C). Mice treated at 6 and 8 weeks were not different from each other (6.8 ± 2.3 weeks, 6.7 ± 0.6  
283 weeks respectively) nor from saline injected mice (6.8 ± 0.7 weeks) (One-way ANOVA with Tukey's  
284 multiple comparisons test). Only mice injected at 4 weeks and *Npc1*<sup>+/+</sup> mice gained weight from 6-9  
285 weeks (11.2% ± 13.1%, 6.0% ± 4.1% respectively), and only mice injected at 4 weeks were significantly  
286 different from all other groups except *Npc1*<sup>+/+</sup>. *Npc1*<sup>m1N</sup> mice injected at 6 or 8 weeks of age lost weight  
287 similar to saline injected *Npc1*<sup>m1N</sup> mice (-12.2% ± 7.7%, -20.0% ± 10.9%, and -14.6% ± 6.8%,  
288 respectively) (Fig 5D, Ordinary one-way ANOVA with Tukey's multiple comparisons test). Longitudinal  
289 weight data further supports that in both male and female cohorts, mice treated earlier maintained weight  
290 and survived longer (Fig S2C, D).

291

292 Evaluating disease progression via the composite phenotype assessment demonstrated benefit early in the  
293 disease course for the group injected at 4 weeks old. Between 6 and 9 weeks of age, mice injected pre-  
294 symptomatically (4 weeks old) had significantly lower phenotype scores than the other two treatment  
295 groups. From 9-12 weeks, all treated groups progressed at a slower rate than saline injected mice, and  
296 mice injected at 4 weeks maintained significantly lower scores than all other groups apart from *Npc1*<sup>+/+</sup>

297 and lived longer (Fig 5E) (two-way ANOVA mixed effects model with Tukey's multiple comparisons  
298 test, results presented in Fig S2B).

299  
300

### 301 **Earlier treatment leads to greater transduction efficiency and reduces cerebellar pathology.**

302

303 Further analyses were performed to determine the differential transduction efficiency of the same vector  
304 at different ages of injection. ddPCR was used to measure *hNPC1* copy number in the cerebrum and liver.  
305 In the cerebrum, for both 4- and 6-week-old treated mice, higher copy numbers predicted longer lifespans  
306 suggesting transduction in the brain aids in lengthening lifespan as shown by linear regression (Fig 6A).  
307 Copy number in the brain did not predict lifespan in mice treated at 8 weeks old. Of note, in the 4-week-  
308 old treated mice, there are two outliers (same from medium-dose group) that do not drive significance.  
309 For further corroboration, copy number was also evaluated across all groups at an age-matched time point  
310 of 9 weeks (late-stage disease, shortly before humane endpoint). At this time point, mice injected post-  
311 symptomatically (8 weeks old) had the highest copy number in the cerebrum and were significantly  
312 different than saline injected and other treated groups (Fig 6B).

313

314 In the brain, immunofluorescent imaging was used to assess neuroinflammation and Purkinje neuron  
315 survival in 9-week-old, age-matched mice. When assessing overall loss of Purkinje neuron survival and  
316 particularly in anterior lobules of the cerebellum, treatment with AAV9 at 4 weeks, as compared to saline-  
317 injected mice and mice treated at 6 or 8 weeks, appears to delay Purkinje neuron loss (Fig 6C). Similarly,  
318 it appears that early treatment at 4 weeks reduces neuroinflammation in posterior lobules of the  
319 cerebellum, as demonstrated by reduced IBA1 staining when compared to saline-injected mice and mice  
320 treated at later time points (Fig 6D).

321

322 In the liver, higher copy numbers predicted increased lifespan in only 6-week-old treated mice; all other  
323 relationships between copy number and age were not predictive in liver (Fig S3A). At 9 weeks of age,  
324 mice treated at 8 weeks again had higher *hNPC1* copy numbers than other treated mice and saline-  
325 injected mice (Fig S3B). At the same age, myeloid cell pathology in showing liver lipid storage burden  
326 reveals reduced CD68+ area in 6-week-old treated mice compared to mice treated at 4 weeks and saline-  
327 injected mice (Fig S3C, D).

328

329

### 330 **AAV9-EF1a(s)-*hNPC1* improved lifespan, delayed disease progression, and decreased liver 331 pathology in *Npc1*<sup>I1061T</sup> mice.**

332

333 To evaluate whether AAV9-EF1a(s)-*hNPC1* was effective in a hypomorphic mouse model of NPC1  
334 disease, *Npc1*<sup>I1061T</sup> mice were treated with 1.28x10<sup>14</sup> gene copies/kg of AAV9-EF1a(s)-*hNPC1* at 4 weeks  
335 old. Survival of treated mice was compared to survival of saline-injected mice (Fig 7A). AAV9 treated  
336 mice lived to a median age of 22.0 weeks which was significantly longer than 15.0 weeks for saline-  
337 injected mice (Log-rank Mantel-Cox test, P<0.001). Of note, *Npc1*<sup>I1061T</sup> mice live to a median survival of  
338 about 17.9 weeks, whereas *Npc1*<sup>m1N</sup> mice have a median survival of about 10.5 weeks [47].

339

340 To evaluate AAV9 efficacy in slowing a common marker of NPC1 progression, weight loss, week of  
341 peak weight and percent weight change from 10 to 14 weeks was measured (adjusted from null model  
342 based on lifespan of *Npc1*<sup>I1061T</sup> model). Treated mice appear to reach peak weight later than saline-injected  
343 mice, but were not significantly different (12.9 ± 3.8 weeks vs. 10.9 ± 1.1 weeks, Wilcoxon matched pairs  
344 signed rank test) (Fig 7B). Treated mice gained weight from 10-14 weeks (3.8% ± 6.0%), similar to  
345 *Npc1*<sup>+/+</sup> mice (7.3% ± 2.7%) (Kruskal-Wallis test with Dunn's multiple comparisons test, P=0.02). In  
346 contrast, saline injected mice lost weight (-12.3% ± 11.0%) and were significantly different from treated  
347 (P<0.05) and *Npc1*<sup>+/+</sup> (P<0.0001) (Fig 7D).

348  
349 Disease phenotype was also evaluated in treated and untreated mice to determine efficacy of gene therapy  
350 in slowing progression. Mice were evaluated on the same metrics as previously described where higher  
351 composite scores indicate worsening disease severity. From weeks 9-15, AAV9 treated *Npc1*<sup>I1061T</sup> mice  
352 had significantly lower composite scores compared to saline-injected mice (9 weeks: P=0.03; 12-15  
353 weeks: P<0.0001; two-way ANOVA with Tukey's multiple comparisons test), though were still  
354 significantly higher than *Npc1*<sup>+/+</sup> from 9-15 weeks (P<0.0001, Fig 7D).

355  
356 To determine the ability of the AAV9 to transduce cells, ddPCR was used to measure *hNPC1* copy  
357 number in the cerebrum and liver. In both tissues, there were significant relationships between *hNPC1*  
358 copy number and age of humane endpoint; however, the relationship was positive in the cerebrum, and  
359 negative in the liver (Fig S4 A, C). At 14 weeks, the typical end stage for untreated mice, copies of  
360 *hNPC1* in both the cerebrum and liver were present, like values observed in treated *Npc1*<sup>m1N</sup> mice (Fig S4  
361 B, D, Fig 2 A, E).

362  
363 To further analyze therapeutic efficacy, liver lipid storage burden was assessed with  
364 immunohistochemical staining of myeloid cells in age-matched 14-week-old mice. *Npc1*<sup>I1061T</sup> mice show  
365 myeloid cell enlargement due to cholesterol accumulation unlike normal livers from *Npc1*<sup>+/+</sup> mice.  
366 Treated mice show partial reduction of myeloid cell enlargement compared to saline-injected mice (Fig  
367 S4E, F) suggesting gene therapy is effective in reducing cholesterol accumulation in the liver, albeit not to  
368 normal *Npc1*<sup>+/+</sup> levels.

## 369 370 371 **Discussion**

372  
373 Here, we detail a highly effective AAV9 gene therapy that can be delivered systemically, over a range of  
374 doses and times in the *Npc1*<sup>m1N</sup> model of NPC1 deficiency. Administration of higher doses of AAV9,  
375 especially in the pre-symptomatic period, significantly increased survival, slowed weight loss, and  
376 lessened disease severity compared to mice receiving lower vector doses or when treated later in life.  
377 Additionally, we define the durability of AAV9-EF1a(s)-*hNPC1* using knock-in *Npc1*<sup>I1061T</sup> mouse model  
378 as evidenced by increasing lifespan and ameliorating disease progression.

379  
380 We examined dose, age at intervention, and type of genetic variation (null vs missense mutation) as key  
381 factors for optimization of systemically delivered AAV9-hNPC1. Previous studies have established  
382 systemic and CNS routes of administration of gene therapy for NPC disease to be effective, and notably  
383 that higher doses of vector delivered intracerebroventricular (ICV) improved neurological and motor  
384 symptoms of *Npc1*<sup>m1N</sup> mice when compared to lower doses [21, 31-35]. These studies, however, did not  
385 look at varying doses when injected systemically at a juvenile time point. In accordance, we show higher  
386 doses of AAV9 injected retro-orbitally improved disease progression as demonstrated by significantly  
387 longer lifespans, lower phenotype scores, weight maintenance, and reduced pathology when compared to  
388 lower doses.

389  
390 Some proof-of-concept NPC1 gene therapy preclinical studies have been conducted in neonatal *Npc1*<sup>m1N</sup>  
391 mice. While many NPC1 individuals are severe and present early in life, even in utero, others have  
392 varying ages of onset. Many exhibit the first neurological sign in their childhood/juvenile years followed  
393 by significant diagnostic delay, and NPC1 is not on the RUSP or on the carrier screening panel; therefore,  
394 we sought to assess therapeutic efficacy of AAV9-EF1a(s)-*hNPC1* in early and post-symptomatic disease  
395 states. We show that survival is correlated with cerebral *hNPC1* gene copy number following earlier  
396 intervention, highlighting the importance of early treatment to ameliorate disease progression. Moreover,  
397 *Npc1*<sup>m1N</sup> mice treated at 4 weeks of age, as compared to those treated at 6 or 8 weeks, demonstrate  
398 significant weight gain during disease course and most importantly, survive longer. Of note, the *Npc1*<sup>m1N</sup>



399 mice treated at 6 weeks do show slight benefit from AAV9 treatment, namely modest increase in survival  
400 when compared to mice treated at 8 weeks or saline-injected mice, suggesting that improvements can still  
401 occur when gene therapy is provided at an early symptomatic time point.

402  
403 The *Npc1*<sup>I1061T</sup> model has not been previously investigated. Consistent with the null model, our data  
404 shows a significant increase in survival, delay of weight loss, improvement in phenotype, and reduction of  
405 pathology following systemic administration of AAV9-EF1a(s)-*hNPC1*. Importantly, *Npc1*<sup>I1061T</sup> mice are  
406 more representative of corresponding human condition as NPC1 p.I1061T encodes a misfolded protein  
407 targeted for ERAD [47, 48]. The success of treatment in both mouse models highlights the immediate  
408 translational potential of this vector, and presents opportunities for further investigation, including  
409 examination of other routes of delivery, ages of administration, efficacy in larger animal models, and even  
410 combination with other therapeutic agents to potentially increase efficacy. Importantly, several clinical  
411 trials using peripherally administered gene therapy for neurological diseases are well underway (SMA1  
412 NCT03461289, NCT03306277, and [17]; MPSIII: NCT02716246, NCT03315182; Canavan:  
413 NCT04498396).

414  
415 In aggregate, our studies demonstrate that systemic delivery of AAV9-EF1a(s)-*hNPC1* can have  
416 significant impact on NPC1 disease phenotype and improve survival in severe and milder mouse models  
417 of the disorder. In addition to mitigation of CNS disease, our data suggests correction in the liver can be  
418 achieved with AAV9-EF1a(s)-*hNPC1*, as evidenced by hepatic transgene expression and reduced liver  
419 pathology in the treated mice. Given the great success using peripherally administered AAV9 gene  
420 therapy to treat SMA1, and the extension to a wide range of other neurological disorders, including  
421 MPSIII and Canavan disease, the preclinical enabling studies presented here should serve to facilitate the  
422 clinical translation of a promising new therapy for individuals with NPC1 deficiency.

423

## 424 **Materials & Methods**

425

### 426 **Vector construction and production**

427 The vector, AAV9-EF1a(s)-*hNPC1* was previously described and produced by the University of  
428 Pennsylvania Vector Core [21].

429

### 430 **Animals**

431 All animal work in these studies was carried out in accordance with the National Institutes of Health  
432 Animal Care and Use Committee approved protocols. Heterozygous (BALB/cNctr-*Npc1*<sup>m1N</sup>/J strain;  
433 Jackson Laboratory strain # 003092) *Npc1*<sup>+m1N</sup> mice were crossed to obtain homozygous mutants  
434 (*Npc1*<sup>m1N/m1N</sup>) and wildtype controls (*Npc1*<sup>+/+</sup>). *Npc1*<sup>I1061T</sup> mice were generated by crossing heterozygous  
435 *Npc1*<sup>+I1061T</sup> mice (B6.129-*Npc1*<sup>m1Dso</sup>/J strain; Jackson Laboratory strain # 027704) to obtain homozygous  
436 mutants and *Npc1*<sup>+/+</sup> controls. We will use single allele notation to indicate homozygosity. Mice were  
437 weighed weekly and then more frequently as the disease progression neared humane endpoint. Mice were  
438 euthanized at a predefined humane endpoint which occurred when two of the following four criteria were  
439 met: weight falling below 70% of peak weight, repeatedly falling to side during movement, dull eyes or  
440 palpebral closure of eyes, or reluctance to move.

441

442 As per the ARRIVE Essential 10, the following information refers to the studies contained herein. All  
443 studies included vehicle or untreated mice (mutant or wildtype, respectively) as control groups. The same  
444 control mice from the *Npc1*<sup>m1N</sup> line were used for the dose and age at injection studies while the  
445 appropriate groups of control mice from the *Npc1*<sup>I1061T</sup> line were used for the corresponding study. A total  
446 of 238 mice were used in these studies. Group sample size is stated in the legend or figure for each  
447 analysis. Data for dose and age of treatment study includes all saline and medium dose AAV9 injected  
448 *Npc1*<sup>m1N</sup> mice as well as untreated *Npc1*<sup>+/+</sup> at 4 weeks. No specific exclusion criteria were set a priori, and

449 all mice included in the studies were randomly assigned to treatment or control groups using a blocking  
450 method. Except for the researcher overseeing the studies, those involved in gene therapy administration  
451 and data acquisition remained blinded to the greatest extent possible. Evaluators were always blinded to  
452 treatment and genotype when performing the phenotype assessment. The primary outcome measure was  
453 survival. Secondary outcome measures included behavioral assessments, weight, gene copy number, and  
454 pathology. Details of statistical analyses are found in either the methods or figure legends. Experimental  
455 animals, procedures, and results are contained with the methods and results of this publication.

456

#### 457 **Phenotypic Assessment**

458 Mice were tested starting at 6 weeks of age and every three weeks thereafter until humane endpoint or  
459 inability to complete the evaluation. The phenotype score evaluates five behaviors associated with the  
460 NPC1 phenotype in diseased mice as previously described: hindlimb clasp, motor function, kyphosis,  
461 grooming, and a balance-ledge test for cerebellar ataxia [49]. Each phenotype is scored from 0 to 3 with  
462 increasing scores representing a more compromised disease state.

463

464 Phenotypic testing was carried out in a blinded fashion such that individual mice within a cage had  
465 distinct tail markings. Evaluators had access only to cage card numbers and tail markings to identify mice.  
466 The order in which mice were tested varied for each testing date. All animals were group housed.

467

#### 468 **Administration of vector**

469 Treated *Npc1<sup>m1N</sup>* mice received a retro-orbital injection of AAV9-EF1a(s)-*hNPC1* at 4 weeks (weaning), 6  
470 weeks, or 8 weeks of age. *Npc1<sup>I1061T</sup>* mice received a retro-orbital injection of AAV9-EF1a(s)-*hNPC1* at 4  
471 weeks old. Control littermate *Npc1<sup>m1N</sup>* or *Npc1<sup>I1061T</sup>* received a retro-orbital injection of 0.9% saline at 4  
472 weeks or the specified age. Mice were anesthetized using isoflurane for 30-60s and then injected retro-  
473 orbitally with a 30-gauge needle affixed to a 0.3 cc syringe. The study compared *Npc1<sup>m1N</sup>* mice  
474 administered vehicle (saline) or gene therapy vector (AAV9-EF1a(s)-*hNPC1*) at different ages and  
475 different doses. Some control *Npc1<sup>+/+</sup>* mice received retro-orbital injections of saline while others  
476 remained un-injected.

477

478 Across cohorts, mice at 4 weeks were given a dose of  $1.2 \times 10^{12}$  gene copies/mouse, which was an average  
479 of  $1.28 \times 10^{14}$  vector genomes/kg (vg/kg, medium dose). For the age at injection study, the weight at  
480 injection was used to calculate the volume necessary to deliver  $1.28 \times 10^{14}$  vg/kg for each mouse and  
481 subsequently administered at either 6 or 8 weeks of age. For the dosage study, the low dose was  $7.87 \times 10^{12}$   
482 vg/kg and the high dose was  $3.06 \times 10^{14}$  vg/kg (maximum dose allowed by vector concentration).

483

#### 484 **Tissue collection and homogenization**

485 Mice were anesthetized with an intraperitoneal injection of Avertin (lethal dose of 0.04 mL/gm) for  
486 euthanasia as previously described [33]. When mice were insensate, the chest cavity was opened, and  
487 mice were perfused with 0.9% saline. Immediately, half of the brain, one lobe of the liver, and a piece of  
488 spleen, kidney, lung, and leg muscle were collected and frozen on dry ice for tissue homogenization. Mice  
489 were then perfused again with 4% paraformaldehyde to fix tissues; remaining organs (half of the brain,  
490 liver) were collected and stored post-fixation in 4% PFA overnight and then rinsed and stored in PBS.

491

492 A Benchmark Scientific Beadbug homogenizer was used to homogenize frozen tissue with UltraPure  
493 water. Tissue was placed in tubes with 3 mm zirconium beads (cerebrum, cerebellum, brainstem, liver,  
494 spleen, leg muscle) or 1.5 mm zirconium beads (kidney, lung) and homogenized 3 times for 30 seconds at  
495 speed 400. Resulting homogenate was aliquoted into tubes for DNA extraction and protein analysis, the  
496 latter of which also had RIPA buffer with proteinase inhibitor cocktail (11 836 170 001; Millipore Sigma,  
497 Burlington, MA) [33].

498

#### 499 **Western blotting**

500 Protein levels from cerebrum, cerebellum, and liver homogenates were quantified using a BCA Assay  
501 (23225, Thermo Fisher Scientific, Waltham, MA). Equal amounts of protein (50 ug for liver, cerebellum,  
502 and cerebrum) were run on 4-12% Bis-Tris SDS-polyacrylamide gels (NP0321BOX/Invitrogen by  
503 Thermo Fisher Scientific) and separation was achieved via electrophoresis; protein was then transferred to  
504 a nitrocellulose membrane (Life Technologies) and blocked for 1 hour in 0.01% PBSTween and 5% BSA.  
505 Samples were incubated for 1 hour at room temperature on a rocker with primary antibody (Table S2).  
506 Secondary antibodies were incubated for 1 hour at room temperature (Table S2). Imaging of bands was  
507 done with the LI-CORE Odyssey Imaging System.

508

### 509 **Histology**

510 Brain and liver tissue from each group were acquired at 9 or 10 weeks of age (*Npc1<sup>mlN</sup>*), 14 weeks of age  
511 (*Npc1<sup>11061T</sup>*), or humane endpoint (both models). Post-fixation tissues were embedded in agarose blocks  
512 (3.5% agarose, 8% sucrose, phosphate buffered saline - PBS) and sectioned parasagittally (30  $\mu$ m) using a  
513 Leica VT1200 S vibratome. Free-floating sections were collected, incubated in 1.6% H<sub>2</sub>O<sub>2</sub> in PBS, then  
514 washed in 0.25% Triton X-100/PBS (PBSt). After blocking for 1 hour at room temperature in  
515 PBSt/normal goat serum, samples were incubated in primary antibodies overnight at 4°C (Table S2).  
516 Samples were washed in PBSt and then incubated with secondary antibodies for 30 minutes at 37°C  
517 (AlexaFluor 488 or 594; Table S2). Filipin staining (Sigma-Aldrich, F9765) was finally performed to  
518 allow visualization of unesterified cholesterol accumulation (0.05mg/mL) with a 20-minute incubation.  
519 ProLong Gold mounting medium alone (Thermo Fisher Scientific, P36934) or with DAPI (Thermo Fisher  
520 Scientific, P36935) was used to coverslip after mounting slides.

521

522 For immunohistochemical staining after primary antibody incubation, slides were incubated in  
523 biotinylated secondary antibody and washed in PBSt. A biotinylated horse radish peroxidase (HRP) was  
524 preincubated with avidin to form Avidin-Biotin Complex (ABC; Vector Laboratories SK-4100) and the  
525 tissues were incubated in ABC (in PBS) for 1 hour. Tissues were then washed in PBS and incubated for  
526 10 minutes in a 3,3'-Diaminobenzidine (DAB; Vector Laboratories PK-4000) solution before mounting  
527 and cover slipping with VectaMount (Vector Laboratories, H-5700).

528

529 Histoserv, Inc. (Germantown, MD) performed paraffin embedding (FFPE tissues). For  
530 immunofluorescent staining, FFPE sections (3  $\mu$ m) were collected and underwent antigen retrieval in a  
531 citrate (pH 6.0, Electron Microscopy Sciences, 62706-10) or Tris-EDTA (pH 9.0, Abcam, AB93684)  
532 buffer. Slides were then incubated in primary antibody diluted in antibody diluent with BSA and  
533 preservative (Thermo Fisher Scientific, 003218) at 37°C for 1 hour, washed in PBS, and incubated in  
534 secondary antibodies for 30 minutes at 37°C (Table S2). Tissues were then coverslipped with ProLong  
535 Gold mounting medium with or without DAPI.

536

### 537 **Image capture and analysis**

538 Fluorescent imaging was performed with an inverted Zeiss AxioScan Z1slide scanner with Zen Blue 3.8  
539 as previously described in [33]. Brightfield images were captured on the same Zeiss AxioScan Z1 slide  
540 scanner. Adobe Photoshop 2023 (v.23.5.0) and 2024 (v.25.1.0) was used to modify all images in a  
541 figure/group identically by resizing and adjusting brightness and/or contrast.

542

### 543 **Quantification of CD68 area**

544 Percent of positive CD68 area relative to total area in liver sections was determined according to methods  
545 previously described [54] using Image-Pro. V11 software (Media Cybernetics, Inc.). Images were  
546 processed using ten regions of interest with total area 900,000  $\mu$ m<sup>2</sup> to determine average percent positive  
547 area.

548

### 549 **Copy number analysis by ddPCR**

550 Vector copy number was measured by droplet digital PCR (ddPCR) as previously described with NPC1  
551 and GAPDH primers (Bio-Rad; [33]). 0.5-50 ng of DNA was used for gene copy number quantification  
552 with brain (cerebrum, cerebellum, or brainstem) and liver homogenates. Additional organs including  
553 spleen, kidney, lung, and leg muscle were assayed for the 10-week-old cohort in the dose study, using  
554 0.5-5 ng of DNA per reaction. Signals for droplets were either positive or negative for hNPC1 and/or  
555 GAPDH as determined using BioRad's QuantaSoft version 1.7.4.0917 software.

556

#### 557 **Mass spectrometry imaging and lipidomics**

558 The fresh frozen half of the brain was sectioned on a CryoStar NX50 Cryostat set to -12°C in preparation  
559 for mass spectrometry imaging. The frozen tissue was divided into 10 µm thick sections, which were then  
560 promptly thaw-mounted onto ITO slides (MIDSCI) and stored at -80°C. Immediately prior to imaging  
561 sections, the slides were removed from the -80°C freezer and washed with ice cold 50 mM ammonium  
562 formate for 20 seconds then dried in vacuo. 9-aminoacridine and 2,5-dihydroxybenzoic acid were chosen  
563 as the matrices for negative and positive mode respectively. One hundred milligrams of solid matrix was  
564 dissolved in 10 mL of 50:50 H<sub>2</sub>O:ACN + 0.2% TFA and filtered using a 0.2 µm syringe filter. Filtered  
565 matrix was applied to the slide using the HTX TM Sprayer.

566

567 Mass spectrometry imaging was performed on a Bruker RapifleX MALDI TOF with a 10 kHz laser set to  
568 60% power, 500 laser shots per pixel, and a step size of 35 µm. The instrument was operated in negative  
569 and positive mode within an m/z range of 200-1800. All data processing including region-of-interest  
570 determination, spatial segmentation, mass spectra extraction and image generation were performed using  
571 Bruker's FlexImaging software. LIPID MAPS and the Human Metabolome Database were used to  
572 annotate and identify lipids according to accurate mass measurements.

573

#### 574 **Statistical analysis**

575 Randomization was achieved with multiple cohorts. Mice within each cohort were included from each  
576 age at injection group or at each dosage. Statistical analysis was performed using GraphPad Prism version  
577 9.5.1 for Windows or Mac. Normality was evaluated for data sets and appropriate parametric or  
578 nonparametric tests were selected for further analysis. Data is presented as mean ± SD. Kaplan Meier  
579 survival curves used Log-Rank Mantel-Cox test to assess significance, with a Bonferroni-correction  
580 applied for P<0.0083 for multiple (six) comparisons. Other statistical tests were as follows: Kruskal-  
581 Wallis test with Dunn's multiple comparison's test, one-way ANOVA with Tukey's multiple  
582 comparisons test, two-way ANOVA with Tukey's multiple comparisons test, linear regression test (all  
583 multiple comparisons test use post hoc Bonferroni's correction). In all figures: \* P<0.05, \*\* P<0.01, \*\*\*  
584 P<0.001, \*\*\*\* P<0.0001.

585

#### 586 **Acknowledgments:**

587 We thank Laura L. Baxter for her support and instruction in statistical analysis as well as Stephen  
588 Wincovitch for imaging and quantification support. We are grateful for the technical assistance in digital  
589 droplet PCR provided by the National Cancer Institute Genomics Core. We sincerely acknowledge the  
590 NIH animal care and veterinary staff for the care of mice used in these studies. We also recognize the  
591 driving force behind this work: individuals with NPC1 disease. Their perseverance in the face of this  
592 debilitating disease inspires and humbles us.

#### 593 **Funding**

594 This work was supported by the Intramural Research Program of the National Human Genome Research  
595 Institute (NHGRI) at the NIH (1ZIAHG000068-16), the *Eunice Kennedy Schriver* National Institute of  
596 Child Health and Human Development (NICHD) at the NIH (ZIAHD008988), grants from the NIH

597 (R01NS114413, R01NS124784), and the Ara Parseghian Medical Research Fund at the University of  
598 Notre Dame. AVM, ALG, TG, AI, CPV, and WJP were supported by the Intramural Research Program at  
599 NHGRI; CDD, KM, FDP were supported by the Intramural Research Program at NICHD. Additional  
600 support for CDD came from the Support Of Accelerated Research for NPC (Hide & Seek Foundation and  
601 Dana's Angels Research Trust). DP-J was supported by the Bridge to Doctoral Program and the  
602 Diversifying Faculty in Illinois Fellowship.

### 603 **Conflict of Interest Statement**

604 CPV and WJP have NIH patents filed on work related to NPC1 genes and the AAV gene therapy  
605 treatment of NPC1 (US Patent Publication Numbers 20180104289, 20210113635).

606

607 **Figure 1: *Npc1<sup>m1N</sup>* mice treated with AAV9-EF1a(s)-*hNPC1* vector show increased survival and**  
608 **delayed disease phenotype progression.**

609 (A) Kaplan-Meier survival curve of mice treated with low, medium, and high dose AAV9 and saline  
610 injected mice. (B) Table of median survival for each treatment group, significance (Mantel-Cox log rank  
611 test) of data from (A). Level of significance is  $P < 0.0083$  based on Bonferroni's correction with 6  
612 comparisons. (C) Week at which mice reached peak weight (Kruskal-Wallis with Dunn's multiple  
613 comparisons test). (D) Percent weight change between 6 and 9 weeks old (One-way ANOVA with  
614 Tukey's multiple comparisons test). For (C, D): saline  $n = 15$ , low dose  $n = 10$ , medium dose  $n = 24$ , high  
615 dose  $n = 8$ ; for (D): *Npc1<sup>+/+</sup>*  $n = 21$ . (E) Composite phenotype scores for each dosage group with  
616 measurements taken every 3 weeks, starting at 6 weeks. Higher scores indicate disease progression  
617 (results presented in Fig S2A) (saline  $n = 14$ , low  $n = 10$ , medium  $n = 13$ , high  $n = 8$ , *Npc1<sup>+/+</sup>*  $n = 21$ ). For  
618 all: \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ . Data presented as mean  $\pm$  SD for C, D, E.  
619

620 **Figure 2: Efficacy of viral transduction in cerebrum and liver varies depends on dose of AAV9.**

621 (A,B) Analysis of *hNPC1* copy number in the cerebrum (A) and liver (B). (Ai, Bi) Copy number of  
622 *hNPC1* in cerebrum (Ai) and liver (Bi) from 10-week-old mice (One-way ANOVA with Tukey's multiple  
623 comparisons test). For Ai, Bi, C: (saline  $n = 6$ , low  $n = 5$ , medium  $n = 6$ , high  $n = 6$ , *Npc1<sup>+/+</sup>*  $n = 9$ ). (Aii,  
624 Bii) Linear regression of *hNPC1* copy number in the cerebrum (Aii) and liver (Bii) in end stage mice. For  
625 Aii (saline  $n = 14$ , low  $n = 9$ , medium  $n = 20$ , high  $n = 6$ , *Npc1<sup>+/+</sup>*  $n = 17$ ). For Bii (saline  $n = 14$ , low  $n =$   
626  $9$ , medium  $n = 18$ , high  $n = 5$ , *Npc1<sup>+/+</sup>*  $n = 17$ ). (Aiii, Biii) NPC1 protein levels were assessed via western  
627 blot in 10-week-old mice to confirm amount of NPC1 protein in the cerebrum (Aiii) and liver (Biii). For  
628 Aiii, Biii (One-way ANOVA with Tukey's multiple comparisons test) (saline  $n = 6$ , low  $n = 5$ , medium  $n =$   
629  $6$ , high  $n = 6$ , *Npc1<sup>+/+</sup>*  $n = 7$ ). (C) Linear regression of copy number in cerebrum across all treated  
630 groups with lifespan (saline  $n = 14$ , treated  $n = 35$ , *Npc1<sup>+/+</sup>*  $n = 17$ ). (D) Gene copy numbers were  
631 measured for various organs at 10 weeks old. For all: \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ .  
632 Data presented as mean  $\pm$  SD for Ai, Bi, Aiii, Biii, D.  
633

634 **Figure 3: Amelioration of pathology is dose dependent for AAV treated mice at 10 weeks old.**

635 (A) Unesterified cholesterol storage marked by filipin labeling in anterior lobules (lobule IV/V) of the  
636 cerebellum (top row). Astrocytes marked by GFAP labeling (middle row). Purkinje neurons are  
637 visualized by Calbindin D labeling (bottom row) (free floating sections). (B) Microgliosis as evidenced  
638 by IBA1 immunostaining in posterior lobules (lobule IX) of the cerebellum (formalin-fixed, paraffin  
639 embedded sections). For (A, B) Scale bar = 1000 microns for panels and 250 microns for sagittal  
640 cerebellar section. (C, D, E, G) Protein levels for GFAP (C), Calbindin (D), CD68 (E), NPC1 (G) were  
641 assessed via western blot for each 10-week-old mice cohort (One-way ANOVA with Tukey's multiple  
642 comparisons test) (saline  $n = 6$ , low  $n = 5$ , medium  $n = 6$ , high  $n = 6$ , *Npc1<sup>+/+</sup>*  $n = 8$ ). (F) Linear regression  
643 of *hNPC1* in cerebellum and survival of mice (saline  $n = 14$ , low  $n = 9$ , medium  $n = 20$ , high  $n = 6$ ,  
644 *Npc1<sup>+/+</sup>*  $n = 17$ ). For all: \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ . Data presented as mean  $\pm$   
645 SD for C, D, E, G.  
646

647 **Figure 4: Reduction in liver pathology is AAV9 dose dependent.**

648 (A) Cholesterol storage (filipin labeling) and CD68+ immunofluorescent staining in the liver (free  
649 floating sections) in 10-week-old mice. Arrows in high dose inset denote groups of cells without  
650 cholesterol storage. Scale bar for panel = 500 microns, insets scale bar = 50 microns. (B) CD68+  
651 immunohistochemical staining of myeloid cells in livers of 10-week-old mice (free floating sections).  
652 Scale bar = 250 microns. (C) Quantification of percent area CD68 labelled in 10-week-old mice (One-  
653 way ANOVA with Tukey's multiple comparison' test) (saline  $n = 5$ , low  $n = 5$ , medium  $n = 6$ , high  $n = 6$ ,  
654 *Npc1<sup>+/+</sup>*  $n = 9$ ). For C: \*  $P < 0.05$ , \*\*  $P < 0.01$ , \*\*\*  $P < 0.001$ , \*\*\*\*  $P < 0.0001$ , data presented as mean  $\pm$  SD  
655 for C.  
656

657 **Figure 5: *Npc1<sup>m1N</sup>* mice treated with medium dose AAV9 at 4 weeks old show increased survival,**  
658 **delayed weight loss, and slower disease progression than mice treated at 6 or 8 weeks old.**

659 (A) Kaplan-Meier survival curve of mice treated with AAV9. (B) Table of median survival for each  
660 treatment group, significance (Mantel-Cox Log-Rank test) of data from (A) with Bonferroni's correction  
661 for 6 comparisons applied ( $P=0.0083$ ). (C) Week at which mice reached peak weight. (D) Percent weight  
662 change between 6 and 9 weeks old. For C, D (saline  $n = 15$ , 4 weeks  $n = 24$ , 6 weeks  $n = 20$ , 8 weeks  $n =$   
663 20) for D (*Npc1<sup>+/+</sup>*  $n = 21$ ) (one-way ANOVA with Tukey's multiple comparisons test). (E) Composite  
664 phenotype scores for each dosage group with measurements taken every 3 weeks, starting at 6 weeks  
665 (results presented in Fig S2B) (saline  $n = 14$ , 4 weeks  $n=13$ , 6 weeks  $n = 20$ , 8 weeks  $n = 20$ , *Npc1<sup>+/+</sup>*  $n =$   
666 21). For all: \*  $P<0.05$ , \*\*  $P<0.01$ , \*\*\*  $P<0.001$ , \*\*\*\*  $P<0.0001$ . Data presented as mean  $\pm$  SD for C, D,  
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669 **Figure 6: Differential transduction efficacy of AAV9 gene therapy vector injected at varying time**  
670 **points in brain leads to differential amelioration of brain pathology.**

671 (A) Linear regression of *hNPC1* copy number in end stage mice. (saline  $n = 14$ , 4-week  $n = 19$ , 6-week  $n =$   
672 20, 8-week  $n = 20$ , *Npc1<sup>+/+</sup>*  $n = 17$ ). (B) ddPCR was used to measure gene copy number of *hNPC1* at 9  
673 weeks old (One-way ANOVA with Tukey's multiple comparisons test) (saline  $n = 6$ , 4-week  $n = 3$ , 6-  
674 week  $n = 4$ , 8-week  $n = 4$ , *Npc1<sup>+/+</sup>*  $n = 4$ ). (C) Purkinje neurons, labeled by Calbindin D (free floating  
675 sections). (D) Microgliosis, demonstrated by IBA1 staining (formalin-fixed, paraffin embedded sections).  
676 For (C, D): Scale bar = 1000 microns, insets scale bar = 250 microns. For B: \*  $P<0.05$ , \*\*  $P<0.01$ , \*\*\*  
677  $P<0.001$ , \*\*\*\*  $P<0.0001$ , data presented as mean  $\pm$  SD.

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679 **Figure 7: *Npc1<sup>I1061T</sup>* mice treated with AAV9 showed increased survival and growth.**

680 (A) Kaplan Meier curve depicts survival of saline injected *Npc1<sup>I1061T</sup>* mice ( $n = 11$ ) and *Npc1<sup>I1061T</sup>* mice  
681 treated with AAV9 at 4 weeks ( $n = 15$ ). (B) Week mice reached peak weight (Wilcoxon test). (C) Percent  
682 weight change from 10 to 14 weeks (Kruskal-Wallis with Dunn's multiple comparisons). (D) Composite  
683 phenotype score for each group measured from 6 to 21 weeks at 3-week intervals. For B, C, D (saline  $n =$   
684 11, treated  $n = 15$ , *Npc1<sup>+/+</sup>*  $n = 15$ ). For all: \*  $P<0.05$ , \*\*  $P<0.01$ , \*\*\*  $P<0.001$ , \*\*\*\*  $P<0.0001$ . Data  
685 presented as mean  $\pm$  SD.

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## SUPPLEMENTAL

**Table S1: Number of mice used in each figure panel.** For dose study and age of treatment, the same mice populations were used for *Npc1<sup>m1N</sup>* injected with saline, *Npc1<sup>m1N</sup>* medium dose treated at 4 weeks, and *Npc1<sup>+/+</sup>*. Includes (n) for supplemental figures.

**Table S2: Antibodies used for immunohistochemical and immunofluorescent staining** along with manufacturer and dilution information.

**S1: Mouse weights.** *Npc1<sup>m1N</sup>* mice treated with AAV9 vector and *Npc1<sup>+/+</sup>* mice weights over lifespan. For the dose study, (A) weights of males averaged with SD, (B) weights of females with SD. For age at injection: (C) males, (D) females. For *Npc1<sup>11061T</sup>* mice: (E) males, (F) females. (n) listed in figure.

**S2: *Npc1<sup>m1N</sup>* mice treated with AAV9 vector and *Npc1<sup>+/+</sup>* mice phenotype score comparison from 6-12 weeks.** (A) Two-way ANOVA results of comparison of various dose treatment groups from 6-9 weeks, and 9-12 weeks. Results presented in table. (B) Two-way ANOVA results of comparison of various age treatment groups from 6-9 weeks, and 9-12 weeks. Results presented in table. For (A, B) Tukey's multiple comparisons test, level of significance is P=0.0083 based on Bonferroni's correction with 6 comparisons.

**S3: Liver pathology and copy number variation of gene therapy treated mice at different ages.**

(A) Linear regression between hNPC1 copy number and lifespan (saline n = 14, 4-week n = 18, 6-week n = 20, 8-week n = 20, *Npc1<sup>+/+</sup>* n = 17). (B) ddPCR was used to measure gene copy number of *hNPC1* at 9 weeks old. (One-way ANOVA with Tukey's multiple comparisons test) (saline n = 6, 4-week n = 4, 6-week n = 4, 8-week n = 4, *Npc1<sup>+/+</sup>* n = 4). (C) CD68+ labeling of Kupffer cells in the liver in 9-week-old mice. Scale bar = 250 microns. (D) Quantification of percent area CD68 labelled in 9-week-old mice (One-way ANOVA with Tukey's multiple comparisons test) (saline n = 6, 4-week n = 8, 6-week n = 4, 8-week n = 4, *Npc1<sup>+/+</sup>* n = 4). For all: \* P<0.05, \*\* P<0.01, \*\*\* P<0.001, \*\*\*\* P<0.0001. Data presented as mean ± SD for B, D.

**S4: Transduction efficacy of AAV9 in *Npc1<sup>11061T</sup>* mice cerebrum and liver tissue.**

(A, C) ddPCR was used to measure gene copy number of *hNPC1* at end stage. Linear regression between *hNPC1* copy number and lifespan in cerebrum (A) or liver tissue (C). For A: (saline n = 11, treated n = 12, *Npc1<sup>+/+</sup>* n = 14), C: (saline n = 11, treated n = 13, *Npc1<sup>+/+</sup>* n = 14). (B, D) ddPCR was used to measure *hNPC1* copy number at 14 weeks. (B) *hNPC1* copy number per cell compared to *Npc1<sup>+/+</sup>* in the cerebrum. (D) *hNPC1* copy numbers in the liver compared to both saline treated mice and *Npc1<sup>+/+</sup>*. For B, D: (One-Way ANOVA with Tukey's multiple comparisons test) (saline n = 4, treated n = 5, *Npc1<sup>+/+</sup>* n = 5). (E) Immunohistochemical staining of liver of *Npc1<sup>+/+</sup>*, treated, and saline-injected 14-week-old mice. Scale bar = 250 microns. (F): Quantification of percent area CD68 labelled in 14-week-old mice (One-way ANOVA with Tukey's multiple comparisons test) (saline n = 5, treated n = 5, *Npc1<sup>+/+</sup>* n = 6). For all: \* P<0.05, \*\* P<0.01, \*\*\* P<0.001, \*\*\*\* P<0.0001. Data presented as mean ± SD.

**S5: Representative western blots: quantification of NPC1 in cerebrum and liver.**

Representative western blots for quantification of NPC1 in cerebrum (A) and liver (B). Quantification of proteins are shown in Fig 2Aiii (cerebrum) and 2Biii (liver). Red outline denotes immunoreactive NPC1 protein. CNV of *hNPC1* provided for each sample below the blots. Source images for cerebrum (C) and liver (D).

**S6. Representative western blots: quantification of GFAP and CD68 in cerebellum.**



743 Representative western blots for quantification of GFAP (A) and CD68 (B) in cerebellum. Quantification  
744 of proteins are shown in Fig 3C and 3E, respectively. Red outlined denotes immunoreactive GFAP  
745 protein. CNV of *hNPC1* provided for each sample below the blots. Source images for cerebellum GFAP  
746 (C) and cerebellum CD68 (D).

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748 **S7. Representative western blots: quantification of calbindin D and NPC1 in cerebellum.**

749 Representative western blot for quantification of calbindin D (A) and NPC1 (B) in cerebellum.  
750 Quantification of proteins are shown in Fig 3D and 3G, respectively. Red outline denotes immunoreactive  
751 NPC1 protein. CNV of *hNPC1* provided for each sample below the blots. Source image for cerebellum  
752 (C).

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754 **S8: Mass spectrometry imaging of *Npc1*<sup>+/+</sup> and *Npc1*<sup>m1N</sup> mice treated with AAV9 gene therapy.**

755 The lipids displayed include A) Ganglioside GM2 (d18:1/18:0), B) Hexosylceramide HexCer 46:4;O3,  
756 and C) Dihydroceramide Cer 32:2;O3.

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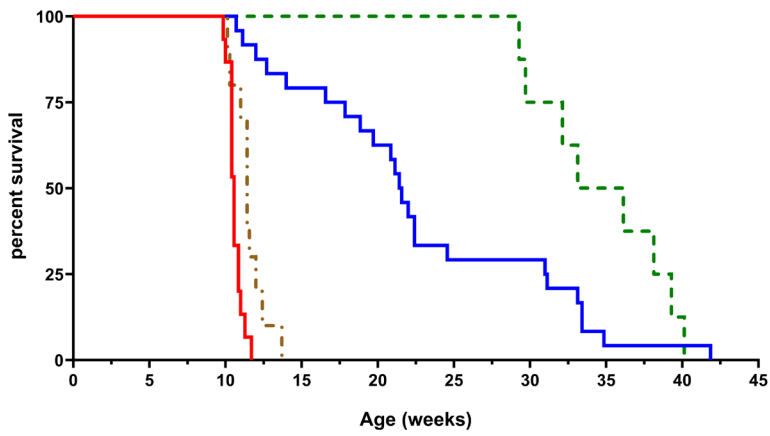
758 **REFERENCES**

- 759 1. Vanier, M.T., *Niemann-Pick diseases*. Handb Clin Neurol, 2013. **113**: p. 1717-21.
- 760 2. Ory, D.S., *Niemann-Pick type C: a disorder of cellular cholesterol trafficking*. Biochim  
761 Biophys Acta, 2000. **1529**(1-3): p. 331-9.
- 762 3. Sleat, D.E., et al., *Genetic evidence for nonredundant functional cooperativity between*  
763 *NPC1 and NPC2 in lipid transport*. Proc Natl Acad Sci U S A, 2004. **101**(16): p. 5886-91.
- 764 4. Infante, R.E., et al., *Purified NPC1 protein. I. Binding of cholesterol and oxysterols to a*  
765 *1278-amino acid membrane protein*. J Biol Chem, 2008. **283**(2): p. 1052-63.
- 766 5. Vanier, M.T., et al., *Genetic heterogeneity in Niemann-Pick C disease: a study using*  
767 *somatic cell hybridization and linkage analysis*. Am J Hum Genet, 1996. **58**(1): p. 118-25.
- 768 6. Cologna, S.M. and A. Rosenhouse-Dantsker, *Insights into the Molecular Mechanisms of*  
769 *Cholesterol Binding to the NPC1 and NPC2 Proteins*. Adv Exp Med Biol, 2019. **1135**: p.  
770 139-160.
- 771 7. Vanier, M.T., *Niemann-Pick disease type C*. Orphanet J Rare Dis, 2010. **5**: p. 16.
- 772 8. Patterson, M.C., et al., *Recommendations for the diagnosis and management of*  
773 *Niemann-Pick disease type C: an update*. Mol Genet Metab, 2012. **106**(3): p. 330-44.
- 774 9. Fields, T., et al., *N-acetyl-L-leucine for Niemann-Pick type C: a multinational double-blind*  
775 *randomized placebo-controlled crossover study*. Trials, 2023. **24**(1): p. 361.
- 776 10. Mengel, E., et al., *Efficacy and safety of arimoclomol in Niemann-Pick disease type C:*  
777 *Results from a double-blind, randomised, placebo-controlled, multinational phase 2/3*  
778 *trial of a novel treatment*. J Inherit Metab Dis, 2021. **44**(6): p. 1463-1480.
- 779 11. Patterson, M.C., et al., *Miglustat for treatment of Niemann-Pick C disease: a randomised*  
780 *controlled study*. Lancet Neurol, 2007. **6**(9): p. 765-72.
- 781 12. Ory, D.S., et al., *Intrathecal 2-hydroxypropyl-beta-cyclodextrin decreases neurological*  
782 *disease progression in Niemann-Pick disease, type C1: a non-randomised, open-label,*  
783 *phase 1-2 trial*. Lancet, 2017. **390**(10104): p. 1758-1768.
- 784 13. Patterson, M.C., et al., *Stable or improved neurological manifestations during miglustat*  
785 *therapy in patients from the international disease registry for Niemann-Pick disease type*  
786 *C: an observational cohort study*. Orphanet J Rare Dis, 2015. **10**: p. 65.
- 787 14. Solomon, B.I., et al., *Association of Miglustat With Swallowing Outcomes in Niemann-*  
788 *Pick Disease, Type C1*. JAMA Neurol, 2020. **77**(12): p. 1564-1568.
- 789 15. Mendell, J.R., et al., *Single-Dose Gene-Replacement Therapy for Spinal Muscular*  
790 *Atrophy*. N Engl J Med, 2017. **377**(18): p. 1713-1722.
- 791 16. Gao, J., R.M. Hussain, and C.Y. Weng, *Voretigene Neparvovec in Retinal Diseases: A*  
792 *Review of the Current Clinical Evidence*. Clin Ophthalmol, 2020. **14**: p. 3855-3869.
- 793 17. Al-Zaidy, S., et al., *Health outcomes in spinal muscular atrophy type 1 following AVXS-*  
794 *101 gene replacement therapy*. Pediatr Pulmonol, 2019. **54**(2): p. 179-185.
- 795 18. Hudry, E. and L.H. Vandenberghe, *Therapeutic AAV Gene Transfer to the Nervous*  
796 *System: A Clinical Reality*. Neuron, 2019. **102**(1): p. 263.
- 797 19. Lowes, L.P., et al., *Impact of Age and Motor Function in a Phase 1/2A Study of Infants*  
798 *With SMA Type 1 Receiving Single-Dose Gene Replacement Therapy*. Pediatr Neurol,  
799 2019. **98**: p. 39-45.
- 800 20. Ali, H.G., et al., *Gene therapy for spinal muscular atrophy: the Qatari experience*. Gene  
801 Ther, 2021. **28**(10-11): p. 676-680.

- 802 21. Chandler, R.J., et al., *Systemic AAV9 gene therapy improves the lifespan of mice with*  
803 *Niemann-Pick disease, type C1*. Hum Mol Genet, 2017. **26**(1): p. 52-64.
- 804 22. Cain, J.T., et al., *Gene Therapy Corrects Brain and Behavioral Pathologies in CLN6-Batten*  
805 *Disease*. Mol Ther, 2019. **27**(10): p. 1836-1847.
- 806 23. Bosch, M.E., et al., *Self-Complementary AAV9 Gene Delivery Partially Corrects Pathology*  
807 *Associated with Juvenile Neuronal Ceroid Lipofuscinosis (CLN3)*. J Neurosci, 2016. **36**(37):  
808 p. 9669-82.
- 809 24. Sondhi, D., et al., *Slowing late infantile Batten disease by direct brain parenchymal*  
810 *administration of a rh.10 adeno-associated virus expressing CLN2*. Sci Transl Med, 2020.  
811 **12**(572).
- 812 25. Manfredsson, F.P., A.C. Rising, and R.J. Mandel, *AAV9: a potential blood-brain barrier*  
813 *buster*. Mol Ther, 2009. **17**(3): p. 403-5.
- 814 26. Foust, K.D., et al., *Intravascular AAV9 preferentially targets neonatal neurons and adult*  
815 *astrocytes*. Nat Biotechnol, 2009. **27**(1): p. 59-65.
- 816 27. Lykken, E.A., et al., *Recent progress and considerations for AAV gene therapies targeting*  
817 *the central nervous system*. J Neurodev Disord, 2018. **10**(1): p. 16.
- 818 28. Verdera, H.C., K. Kuranda, and F. Mingozzi, *AAV Vector Immunogenicity in Humans: A*  
819 *Long Journey to Successful Gene Transfer*. Mol Ther, 2020. **28**(3): p. 723-746.
- 820 29. Bradbury, A.M., et al., *Krabbe disease successfully treated via monotherapy of*  
821 *intrathecal gene therapy*. J Clin Invest, 2020. **130**(9): p. 4906-4920.
- 822 30. Bagel, J.H., et al., *Electrodiagnostic testing and histopathologic changes confirm*  
823 *peripheral nervous system myelin abnormalities in the feline model of niemann-pick*  
824 *disease type C*. J Neuropathol Exp Neurol, 2013. **72**(3): p. 256-62.
- 825 31. Xie, C., et al., *AAV9-NPC1 significantly ameliorates Purkinje cell death and behavioral*  
826 *abnormalities in mouse NPC disease*. J Lipid Res, 2017. **58**(3): p. 512-518.
- 827 32. Hughes, M.P., et al., *AAV9 intracerebroventricular gene therapy improves lifespan,*  
828 *locomotor function and pathology in a mouse model of Niemann-Pick type C1 disease*.  
829 Hum Mol Genet, 2018. **27**(17): p. 3079-3098.
- 830 33. Davidson, C.D., et al., *Improved systemic AAV gene therapy with a neurotrophic capsid in*  
831 *Niemann-Pick disease type C1 mice*. Life Sci Alliance, 2021. **4**(10).
- 832 34. Kurokawa, Y., et al., *Gene Therapy in a Mouse Model of Niemann-Pick Disease Type C1*.  
833 Hum Gene Ther, 2021. **32**(11-12): p. 589-598.
- 834 35. Hughes, M.P., et al., *A Novel Small NPC1 Promoter Enhances AAV-Mediated Gene*  
835 *Therapy in Mouse Models of Niemann-Pick Type C1 Disease*. Cells, 2023. **12**(12).
- 836 36. Tai, C.H., et al., *Long-term efficacy and safety of eladocagene exuparvovec in patients*  
837 *with AADC deficiency*. Mol Ther, 2022. **30**(2): p. 509-518.
- 838 37. Wasserstein, M.P., et al., *The future of newborn screening for lysosomal disorders*.  
839 Neurosci Lett, 2021. **760**: p. 136080.
- 840 38. Patterson, M.C., et al., *Disease and patient characteristics in NP-C patients: findings from*  
841 *an international disease registry*. Orphanet J Rare Dis, 2013. **8**: p. 12.
- 842 39. Surmeli-Onay, O., et al., *Prenatal-onset Niemann-Pick type C disease with nonimmune*  
843 *hydrops fetalis*. Pediatr Neonatol, 2013. **54**(5): p. 344-7.
- 844 40. Kelly, D.A., et al., *Niemann-Pick disease type C: diagnosis and outcome in children, with*  
845 *particular reference to liver disease*. J Pediatr, 1993. **123**(2): p. 242-7.

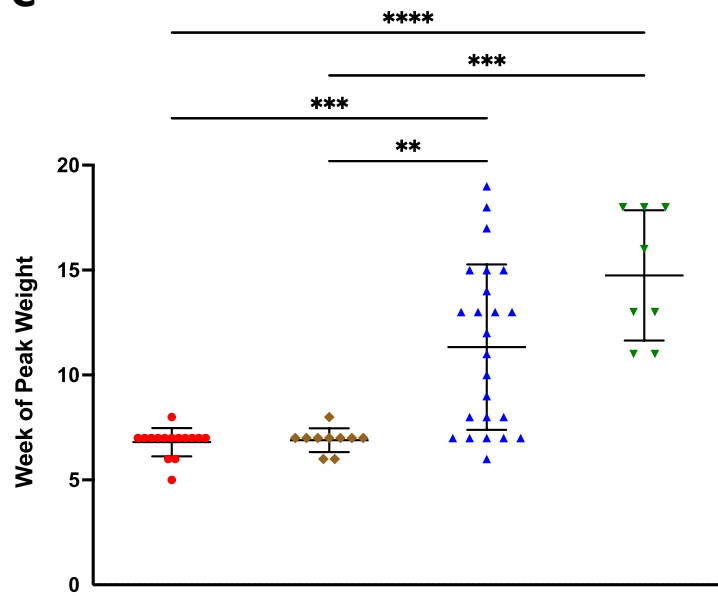
- 846 41. Kumagai, T., et al., *A case of Niemann-Pick disease type C with neonatal liver failure*  
847 *initially diagnosed as neonatal hemochromatosis*. *Brain Dev*, 2019. **41**(5): p. 460-464.
- 848 42. Shammas, H., et al., *Different Niemann-Pick C1 Genotypes Generate Protein Phenotypes*  
849 *that Vary in their Intracellular Processing, Trafficking and Localization*. *Sci Rep*, 2019.  
850 **9**(1): p. 5292.
- 851 43. Landrum, M.J., et al., *ClinVar: public archive of relationships among sequence variation*  
852 *and human phenotype*. *Nucleic Acids Res*, 2014. **42**(Database issue): p. D980-5.
- 853 44. McKay Bounford, K. and P. Gissen, *Genetic and laboratory diagnostic approach in*  
854 *Niemann Pick disease type C*. *J Neurol*, 2014. **261 Suppl 2**(Suppl 2): p. S569-75.
- 855 45. Millat, G., et al., *Niemann-Pick C1 disease: the I1061T substitution is a frequent mutant*  
856 *allele in patients of Western European descent and correlates with a classic juvenile*  
857 *phenotype*. *Am J Hum Genet*, 1999. **65**(5): p. 1321-9.
- 858 46. Guatibonza Moreno, P., et al., *At a glance: the largest Niemann-Pick type C1 cohort with*  
859 *602 patients diagnosed over 15 years*. *Eur J Hum Genet*, 2023. **31**(10): p. 1108-1116.
- 860 47. Praggastis, M., et al., *A murine Niemann-Pick C1 I1061T knock-in model recapitulates the*  
861 *pathological features of the most prevalent human disease allele*. *J Neurosci*, 2015.  
862 **35**(21): p. 8091-106.
- 863 48. Gelsthorpe, M.E., et al., *Niemann-Pick type C1 I1061T mutant encodes a functional*  
864 *protein that is selected for endoplasmic reticulum-associated degradation due to protein*  
865 *misfolding*. *J Biol Chem*, 2008. **283**(13): p. 8229-36.
- 866 49. Yerger, J., et al., *Phenotype assessment for neurodegenerative murine models with*  
867 *ataxia and application to Niemann-Pick disease, type C1*. *Biol Open*, 2022. **11**(4).
- 868 50. Martin, K.B., et al., *Identification of Novel Pathways Associated with Patterned*  
869 *Cerebellar Purkinje Neuron Degeneration in Niemann-Pick Disease, Type C1*. *Int J Mol Sci*,  
870 2019. **21**(1).
- 871 51. Boenzi, S., et al., *Comprehensive-targeted lipidomic analysis in Niemann-Pick C disease*.  
872 *Mol Genet Metab*, 2021. **134**(4): p. 337-343.
- 873 52. Pergande, M.R., et al., *Lipidomic Analysis Reveals Altered Fatty Acid Metabolism in the*  
874 *Liver of the Symptomatic Niemann-Pick, Type C1 Mouse Model*. *Proteomics*, 2019.  
875 **19**(18): p. e1800285.
- 876 53. Zervas, M., K. Dobrenis, and S.U. Walkley, *Neurons in Niemann-Pick disease type C*  
877 *accumulate gangliosides as well as unesterified cholesterol and undergo dendritic and*  
878 *axonal alterations*. *J Neuropathol Exp Neurol*, 2001. **60**(1): p. 49-64.
- 879 54. Rodriguez-Gil, J.L., et al., *Genetic background modifies phenotypic severity and longevity*  
880 *in a mouse model of Niemann-Pick disease type C1*. *Dis Model Mech*, 2020. **13**(3).  
881

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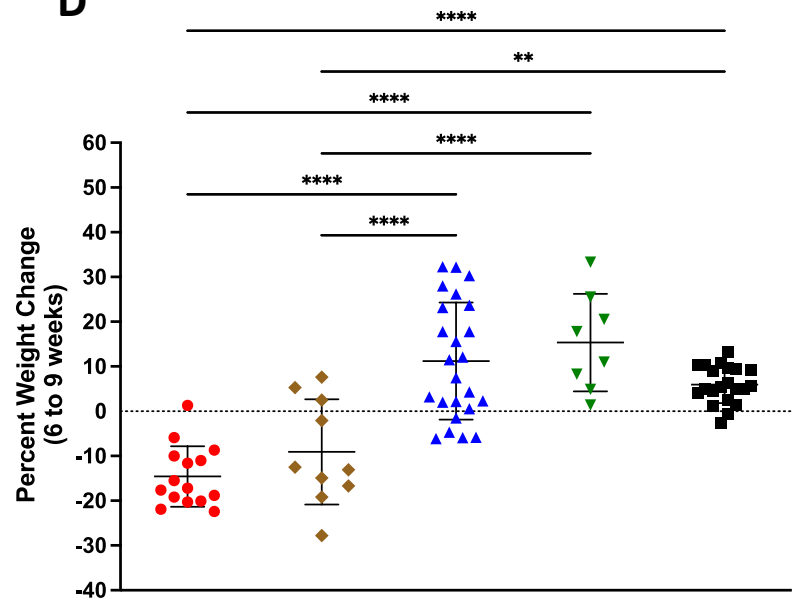


Treatment	Sample Size (n)	Median Survival (weeks)	Significance (Log-rank test)
<i>Npc1<sup>m1N</sup> Saline</i>	15	10.6	
<i>Npc1<sup>m1N</sup> Low</i> 7.87x10 <sup>12</sup> vg/kg	10	11.4	vs. Saline, <i>P</i> =0.0049
<i>Npc1<sup>m1N</sup> Medium</i> 1.28x10 <sup>14</sup> vg/kg	24	21.5	vs. Saline, <i>P</i> <0.0001 vs. Low, <i>P</i> <0.0001
<i>Npc1<sup>m1N</sup> High</i> 3.06x10 <sup>14</sup> vg/kg	8	34.6	vs. Saline, <i>P</i> <0.0001 vs. Low, <i>P</i> <0.0001 vs. Medium, <i>P</i> =0.0266

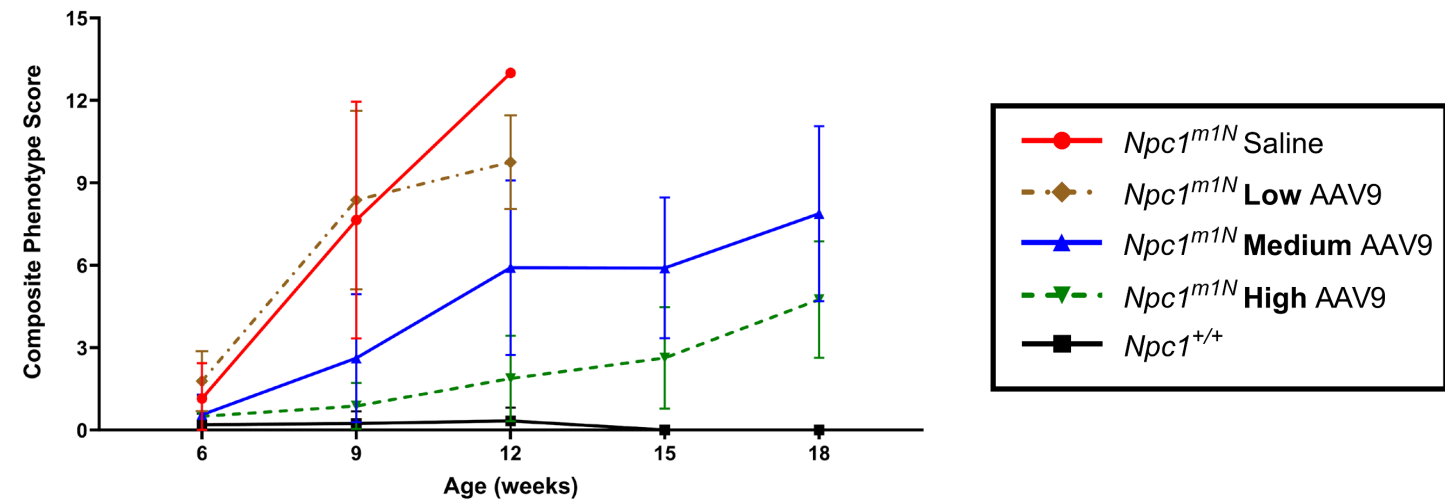
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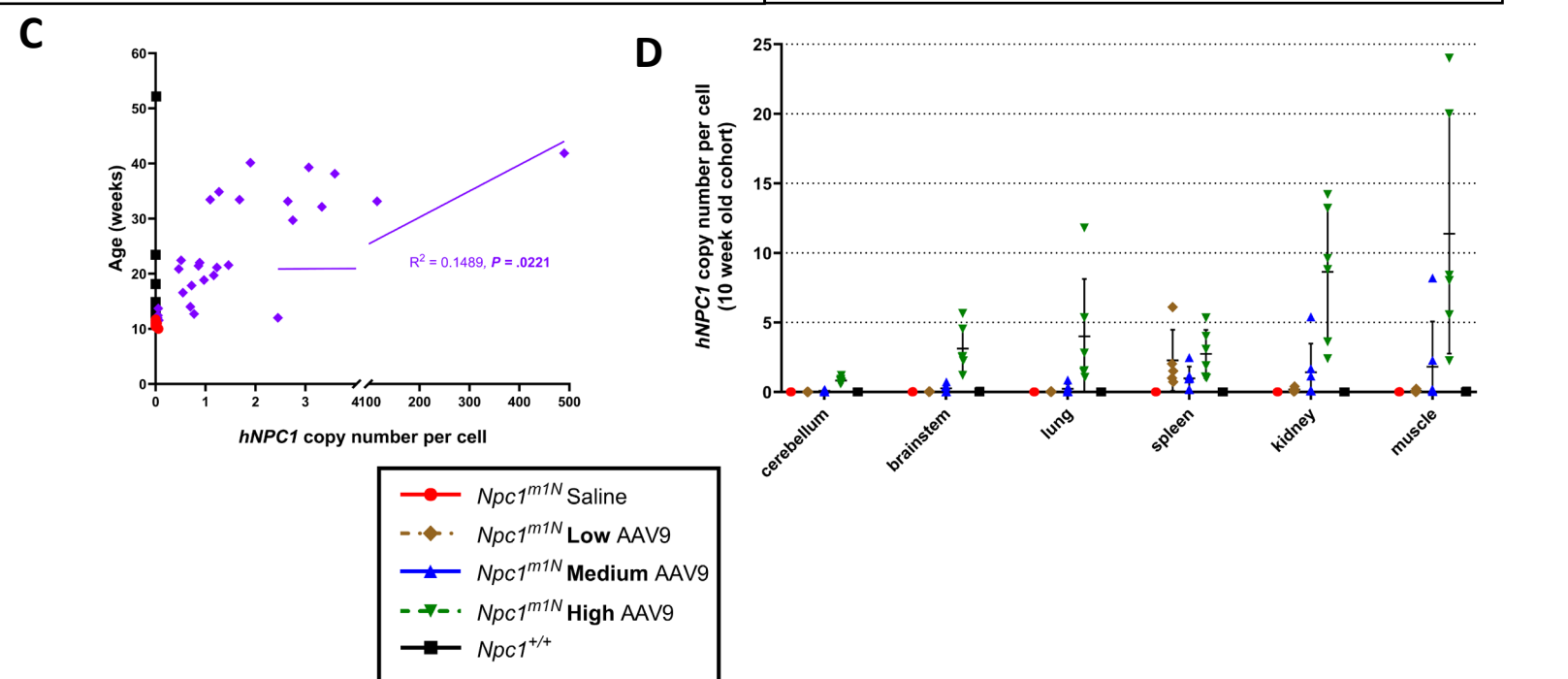
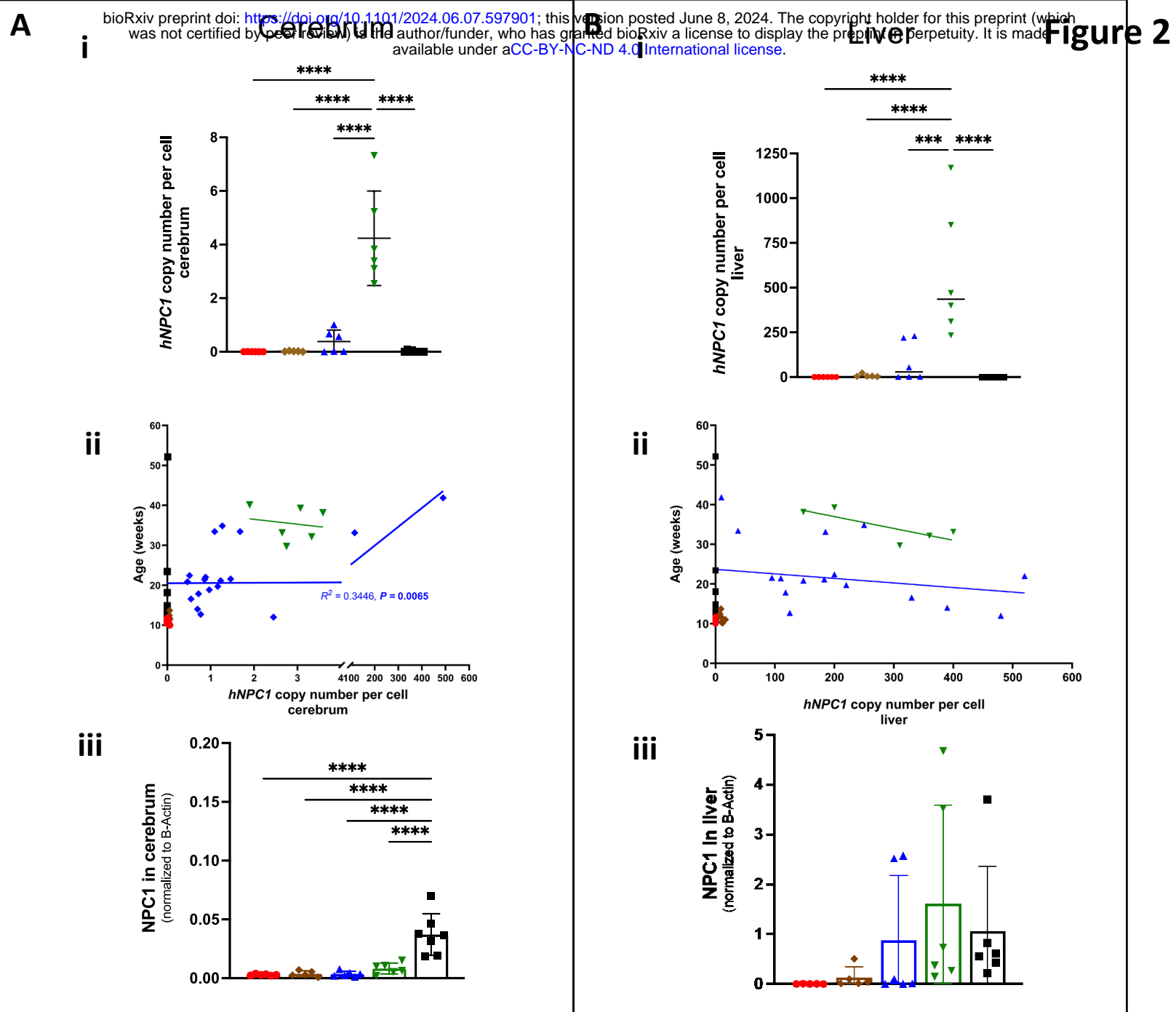


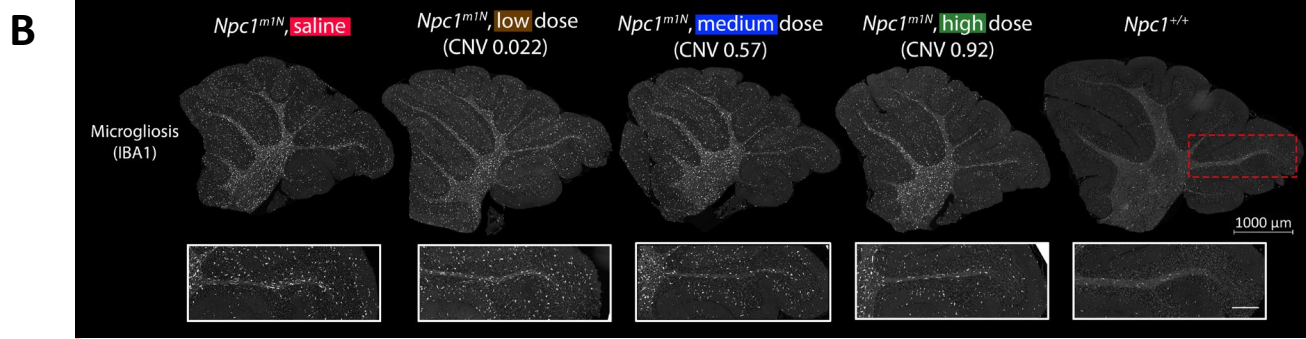
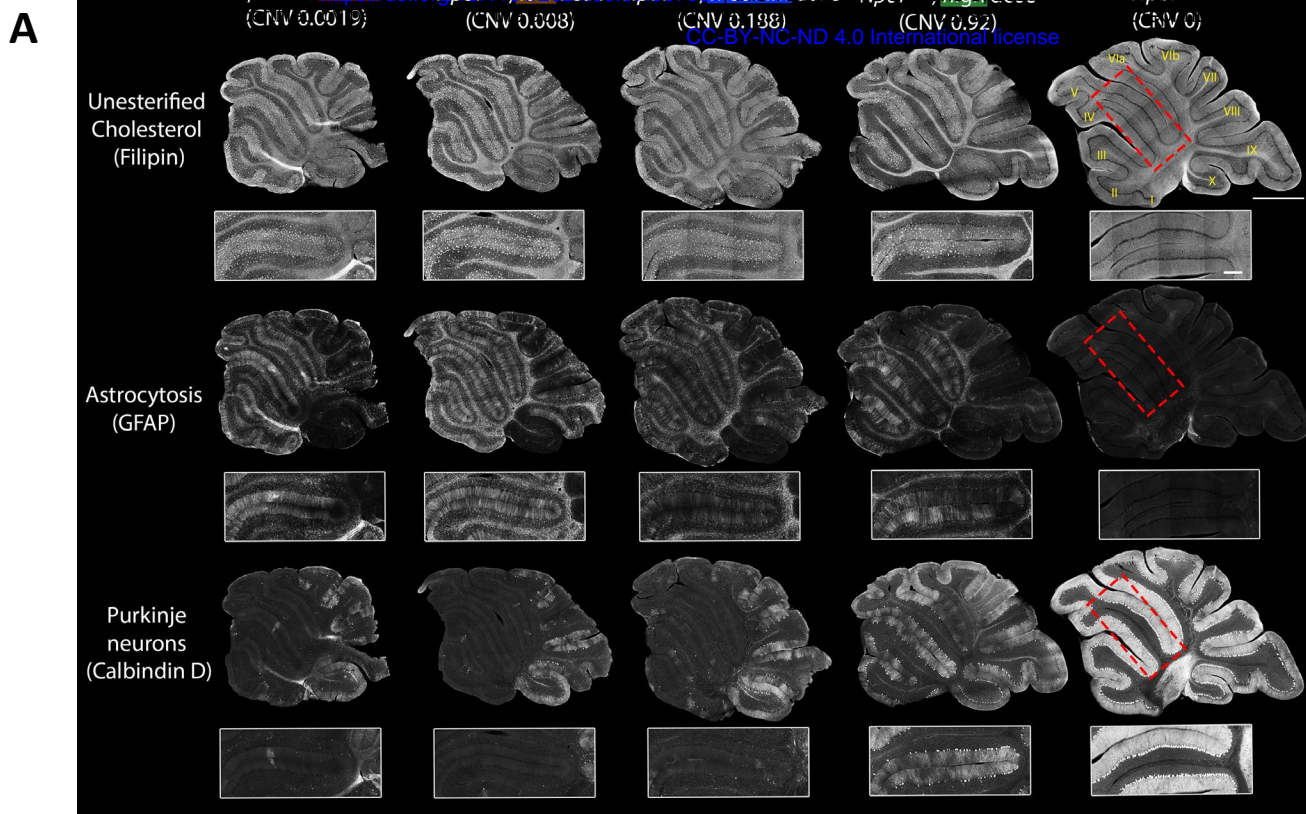
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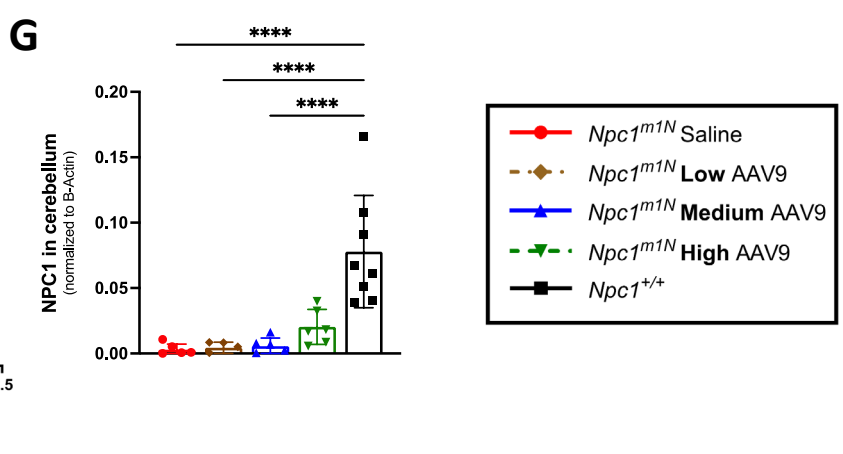
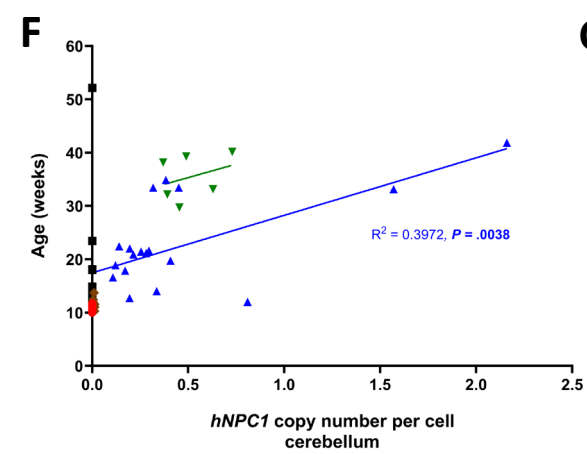
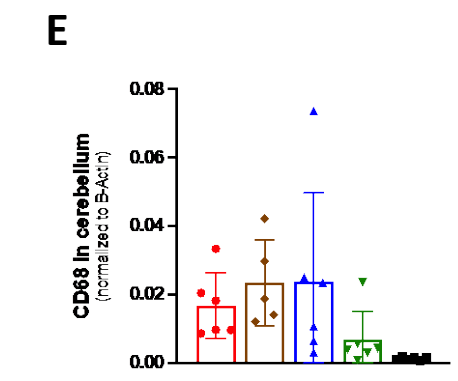
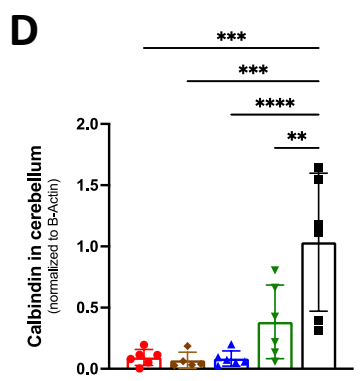
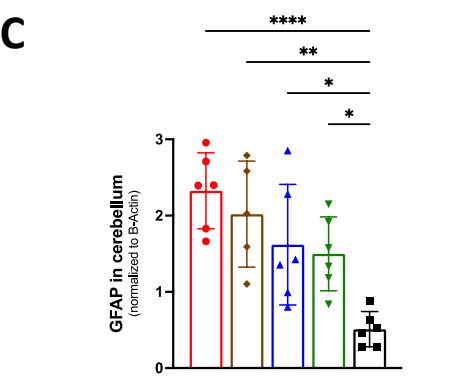
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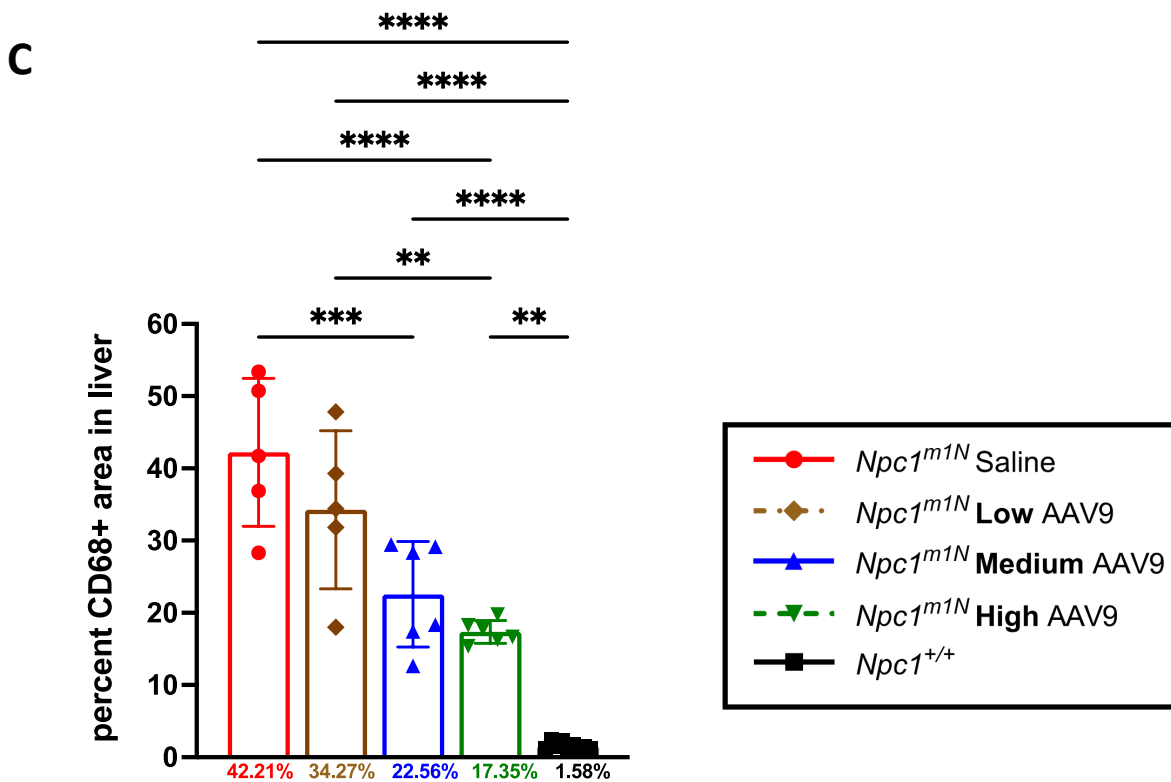
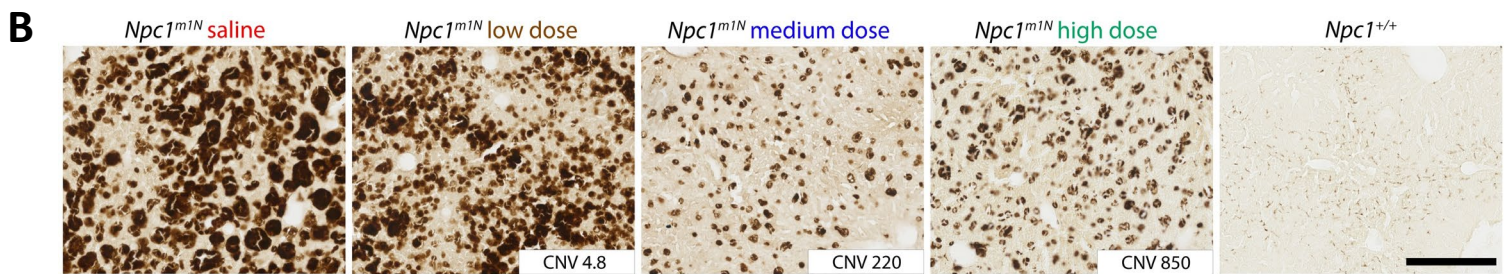
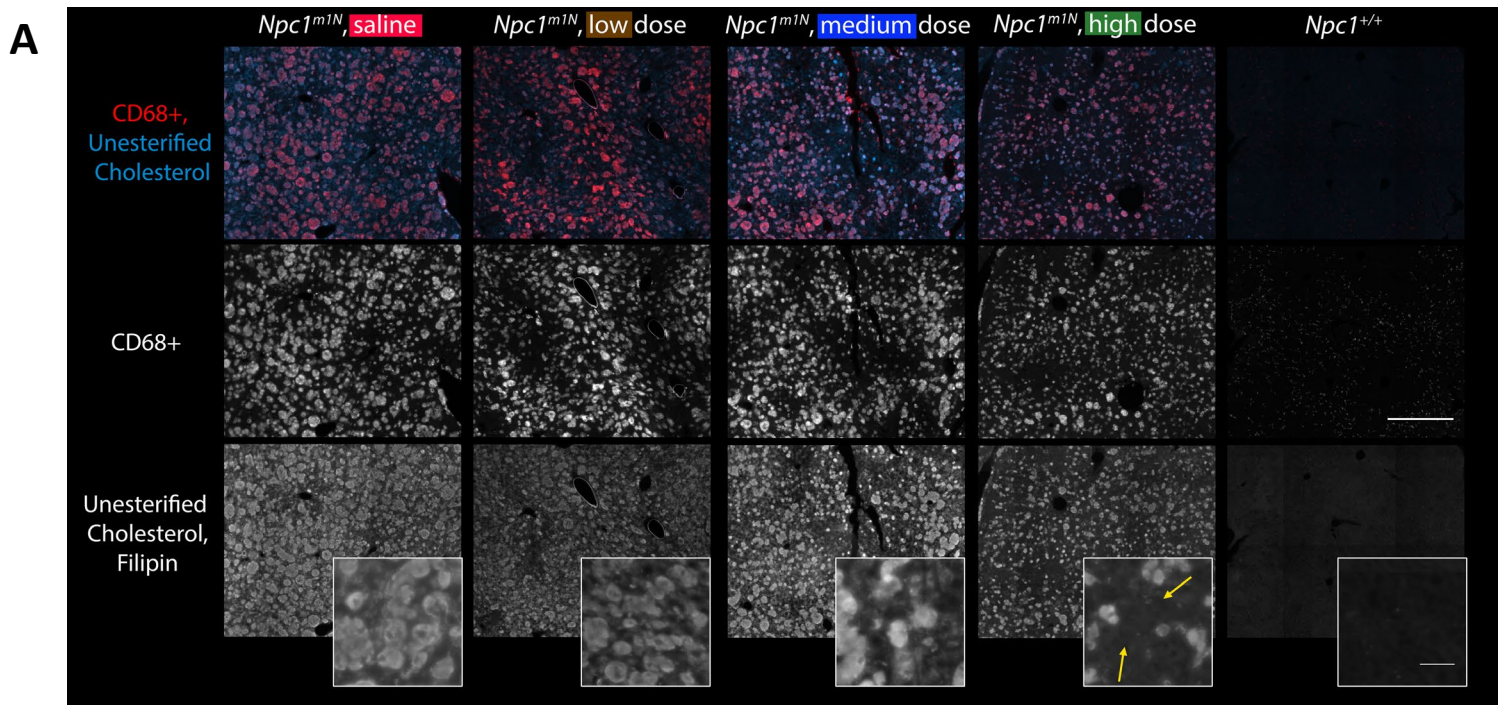




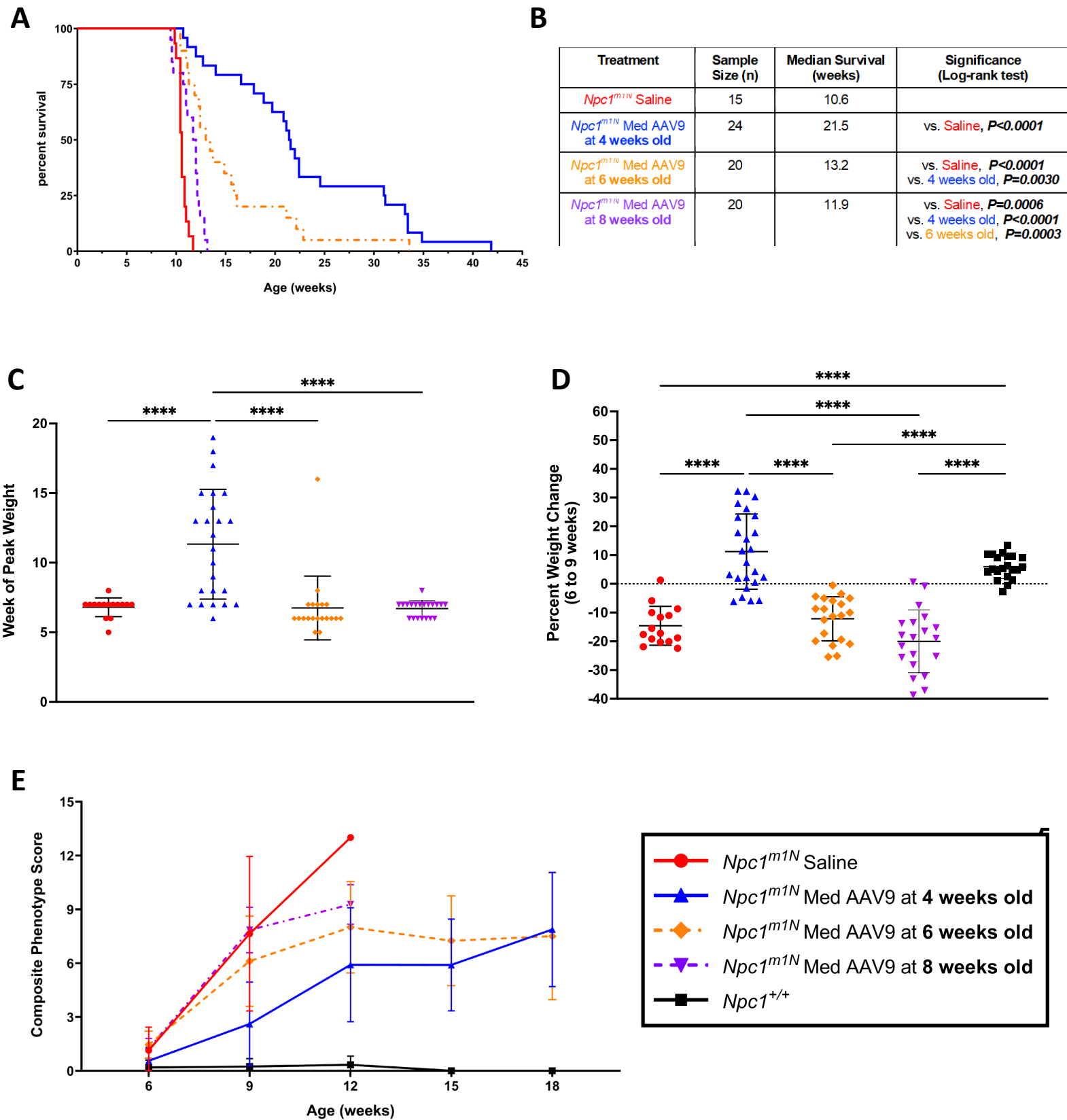


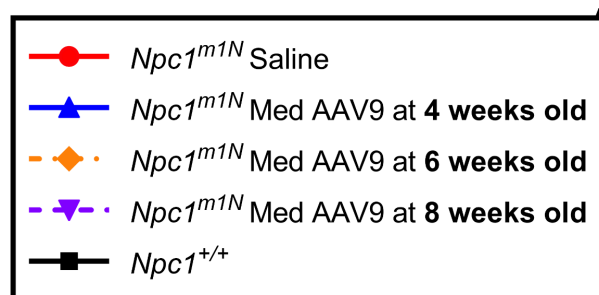
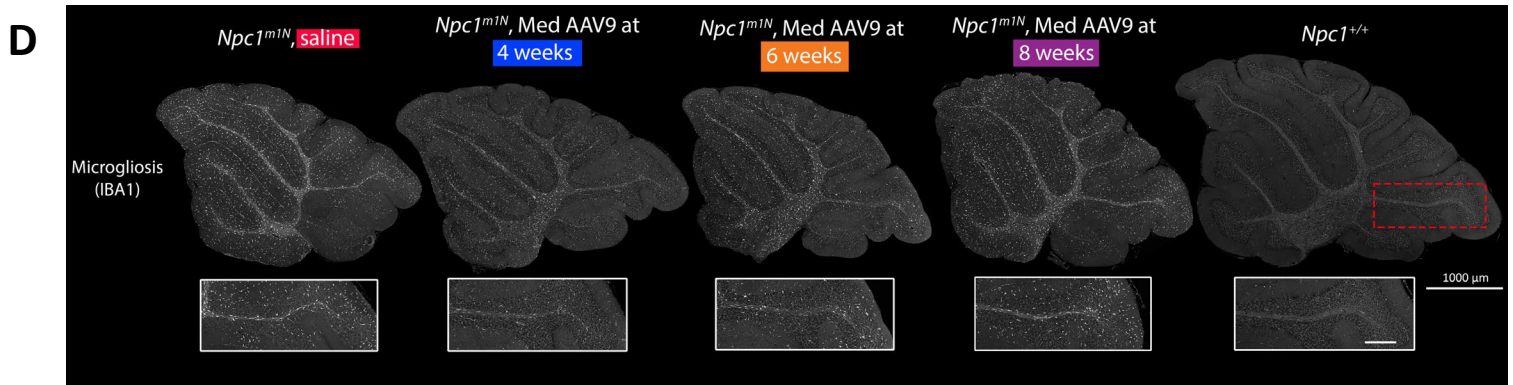
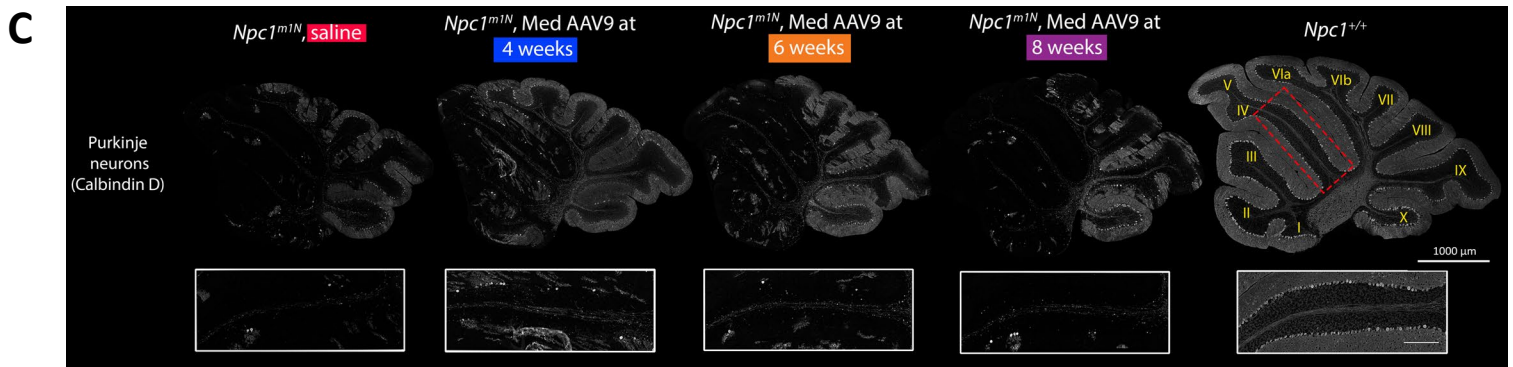
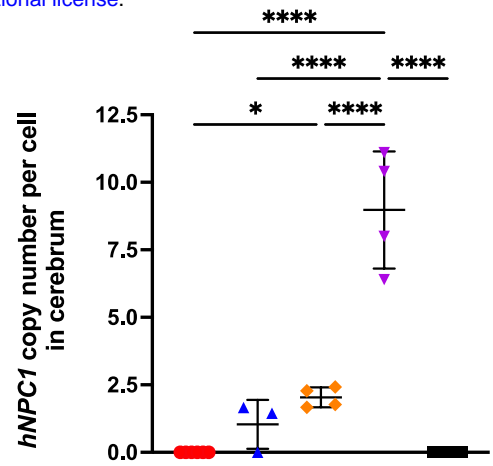
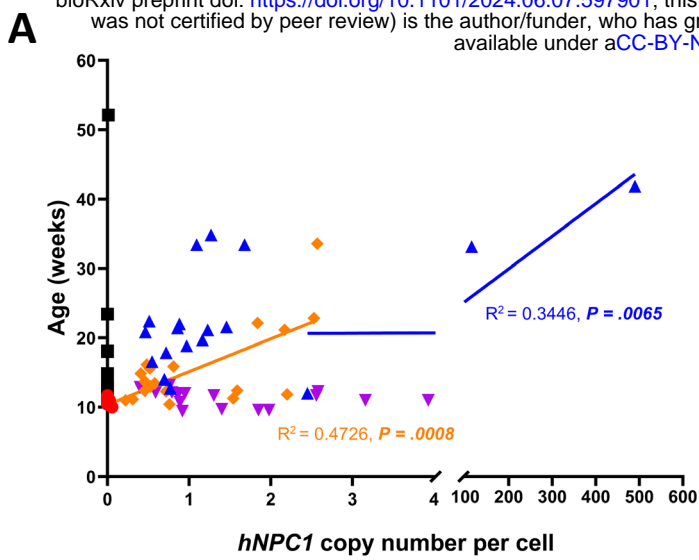
Decreasing pathology











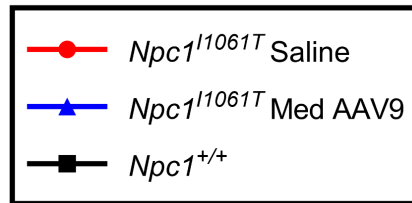
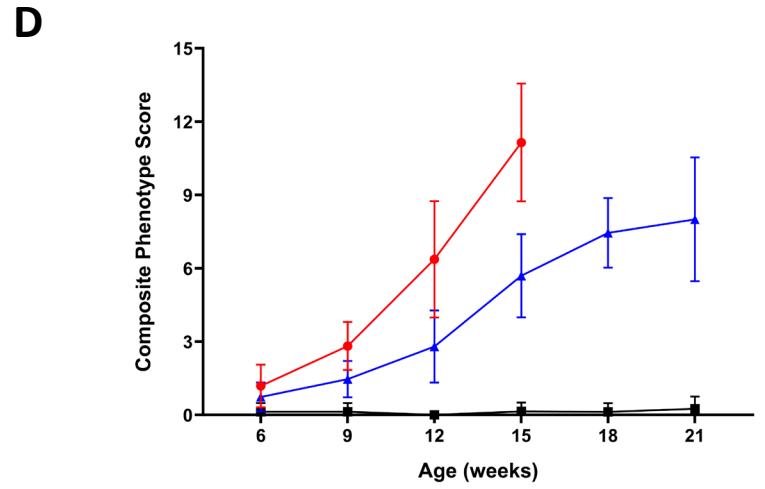
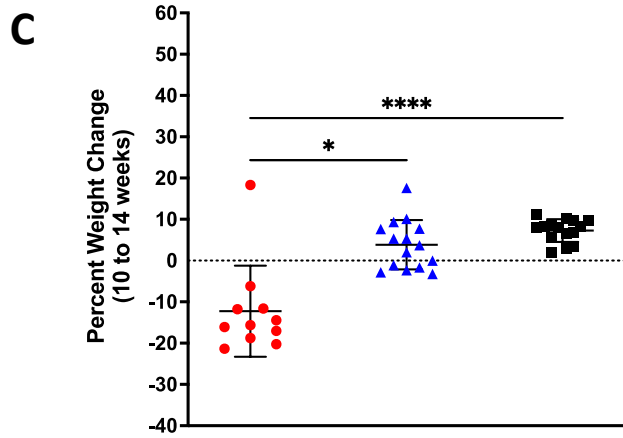
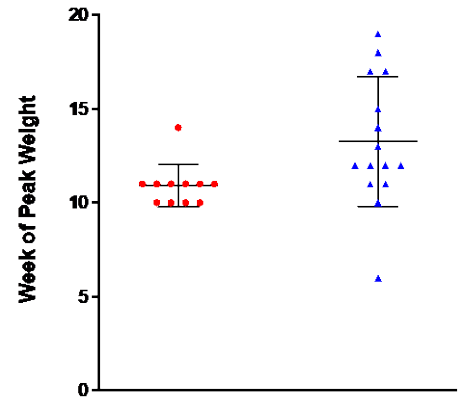
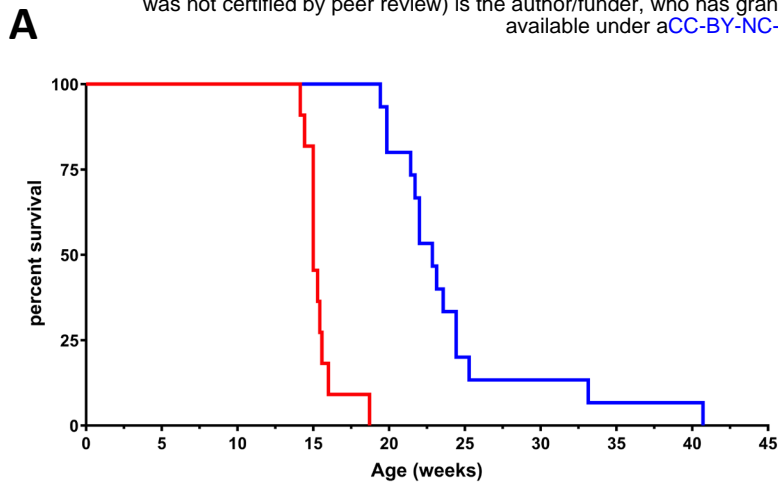
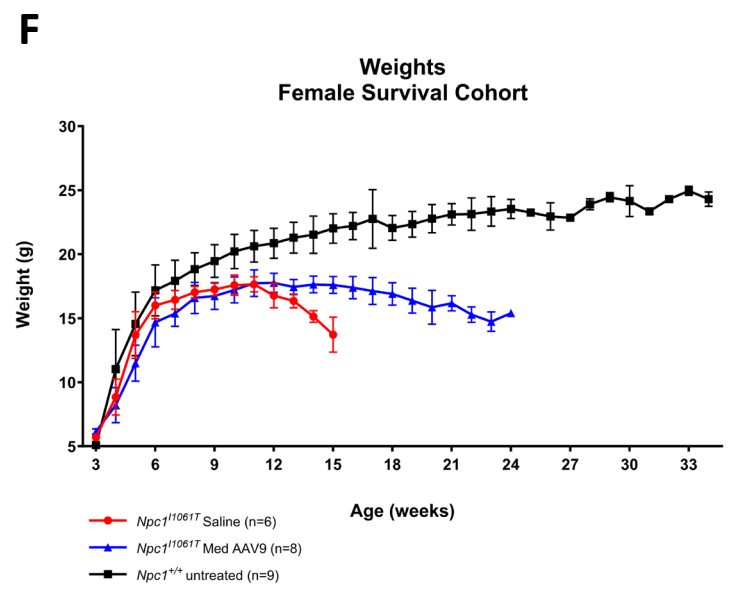
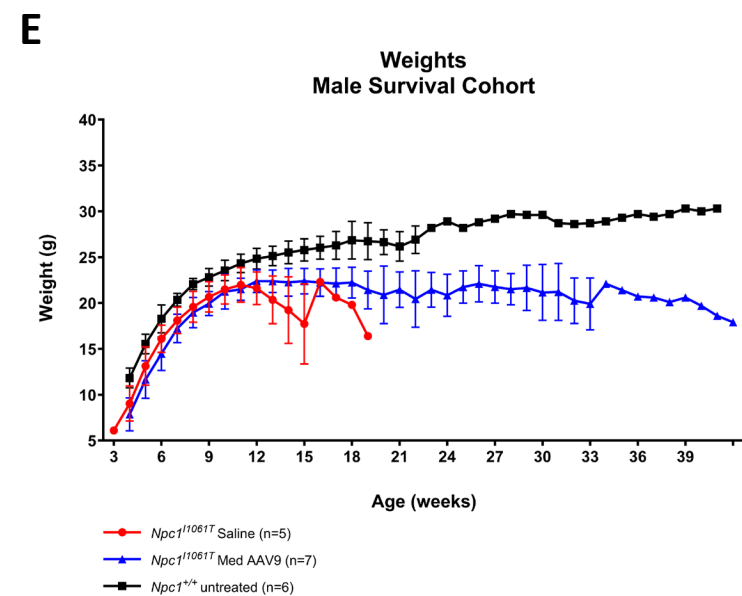
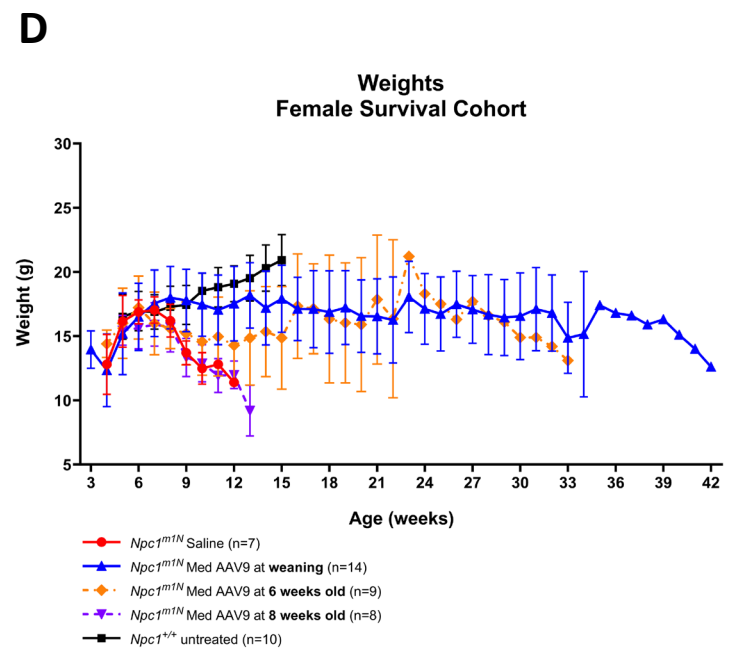
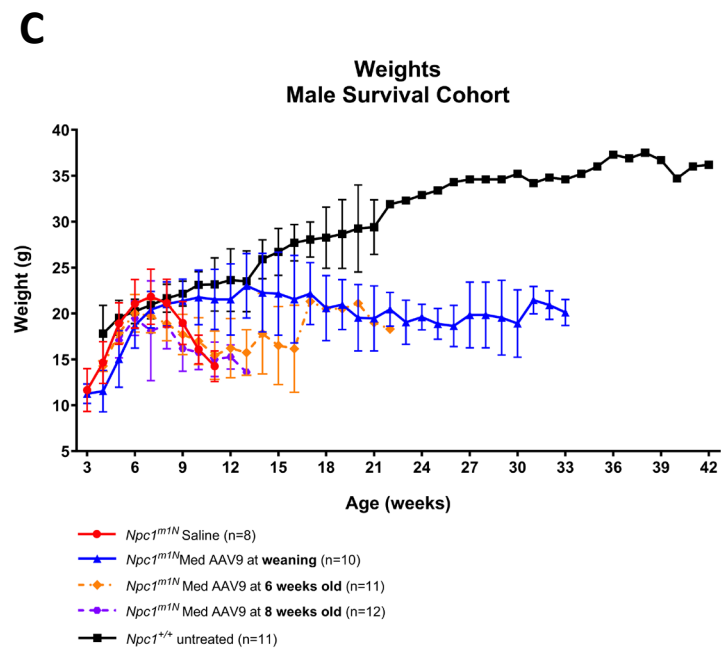
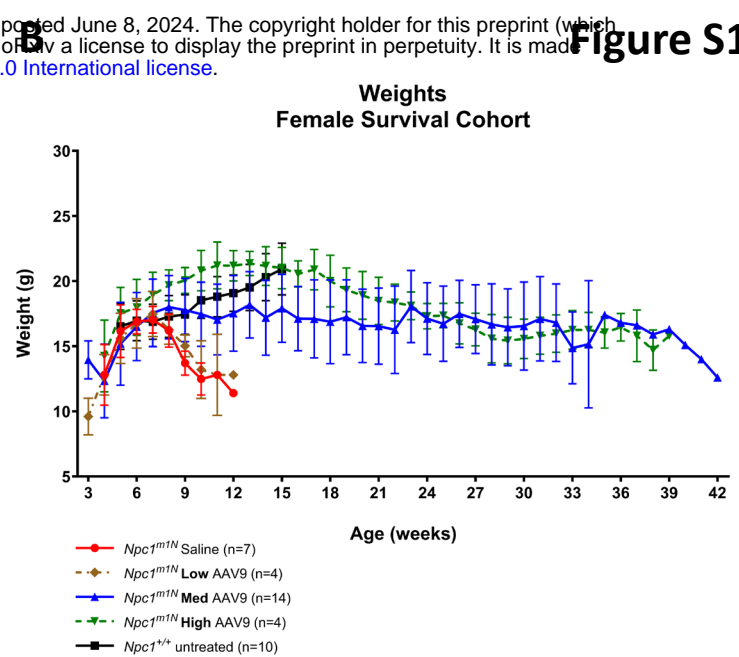
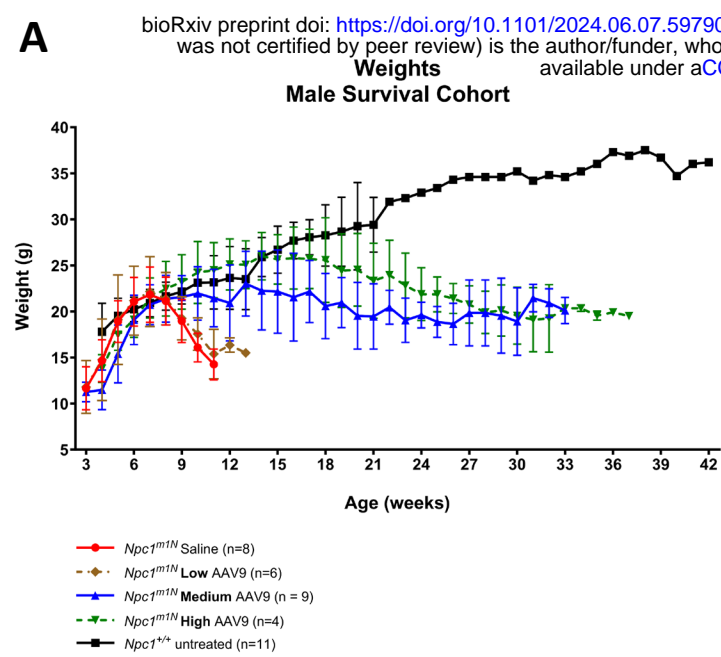


Figure #		<b>Npc1<sup>m1N</sup>, saline</b>	<b>Npc1<sup>m1N</sup>, low</b>	<b>Npc1<sup>m1N</sup>, medium</b>	<b>Npc1<sup>m1N</sup>, high</b>	<b>Npc1<sup>+/-</sup></b>
<b>1</b>	<b>C</b>	15	10	24	8	
	<b>D</b>	15	10	24	8	21
<b>2</b>	<b>A</b>	14	9	20	6	17
	<b>B</b>	6	5	6	5	9
	<b>C</b>	5	5	6	6	7
	<b>D</b>	14	9	18	5	17
	<b>E</b>	6	5	6	6	9
	<b>F</b>	5	5	6	6	6
	<b>G</b>	8		34		16
	<b>H</b>	6	5	6	6	9
<b>3</b>	<b>C</b>	6	5	6	6	8
	<b>D</b>	6	5	6	6	8
	<b>E</b>	6	5	6	6	8
	<b>F</b>	14	9	20	6	17
	<b>G</b>	5	5	6	6	7
	<b>4</b>	<b>C</b>	5	5	6	6
<b>S1</b>	<b>A</b>	8	6	10	4	11
	<b>B</b>	7	4	14	4	10
		<b>Npc1<sup>m1N</sup>, saline</b>	<b>Npc1<sup>m1N</sup>, 4 weeks</b>	<b>Npc1<sup>m1N</sup>, 6 weeks</b>	<b>Npc1<sup>m1N</sup>, 8 weeks</b>	<b>Npc1<sup>+/-</sup></b>
<b>5</b>	<b>C</b>	15	24	20	20	
	<b>D</b>	15	24	20	20	21
<b>6</b>	<b>A</b>	14	20	20	20	17
	<b>B</b>	6	3	3	4	4
<b>S1</b>	<b>C</b>	8	10	11	12	11
	<b>D</b>	7	14	9	8	10
<b>S3</b>	<b>A</b>	14	18	20	20	17
	<b>B</b>	6	4	4	4	4
	<b>D</b>	6	7	4	4	4
		<b>Npc1<sup>I1061T/I1061T</sup>, saline</b>	<b>Npc1<sup>I1061T/I1061T</sup>, medium</b>	<b>Npc1<sup>+/-</sup></b>		
<b>7</b>	<b>C</b>	11	15			
	<b>D</b>	11	15	25		
	<b>E</b>	11	15	15		
	<b>F</b>	5	5	6		
<b>S4</b>	<b>A</b>	11	12	14		
	<b>B</b>	7	5	6		
	<b>C</b>	11	13	14		
	<b>D</b>	7	5	6		

<b>Antibody</b>	<b>Company</b>	<b>Catalog Number</b>	<b>Dilution</b>	<b>Method</b>
<b>α-β-actin (mouse IgG)</b>	ThermoFisher Invitrogen	15G5A11/E2	1:10000	Westerns
<b>α-Calbindin (mouse IgG)</b>	Sigma	C9848	1:750 1:1000	IF Westerns
<b>α-Calbindin (rabbit IgG)</b>	Abcam	ab229915	1:750 (fixed, free-floating sections and FFPE sections)	IF
<b>α-CD68 (rat IgG2a)</b>	BioRad	MCA1957	1:500	IF
<b>α-CD68 (rabbit IgG)</b>	Abcam	ab125212	1:1000	IHC, IF, Westerns
<b>α-GFAP (mouse IgG)</b>	Sigma	G3893	1:1000 (fixed, free-floating sections and FFPE sections) 1:2000	IF Westerns
<b>α-IBA1 (rabbit IgG)</b>	Wako Chemicals	019-19741	1:500 (fixed, free-floating sections) or 1:750 (FFPE)	IF
<b>α-NPC1 (monoclonal rabbit IgG)</b>	Abcam	ab134113	1:2000	Westerns
<b>α-β-III-Tubulin</b>	R&D Systems	MAB1195	1:3000	Westerns
<b>Donkey anti-mouse IgG IRDye 680RD</b>	LI-CORbio	926-68072	1:20000	Westerns
<b>Donkey anti-mouse IgG IRDye 800CW</b>	LI-CORbio	926-32213	1:20000	Westerns
<b>Goat anti-rabbit IgG Biotinylated</b>	Vector Laboratories	BA-1000	1:300	IHC
<b>Goat anti-mouse IgG AlexaFluor 488 or 594</b>	ThermoFisher Invitrogen	A11029 (488) or A11005 (594)	1:350	IF
<b>Goat anti-rabbit IgG AlexaFluor 488 or 594</b>	ThermoFisher Invitrogen	A11034 (488) or A11037 (594)	1:350	IF
<b>Goat anti-rat IgG Alexa Fluor 488 or 594</b>	ThermoFisher Invitrogen	A11006 (488) or A11007 (594)	1:350	IF

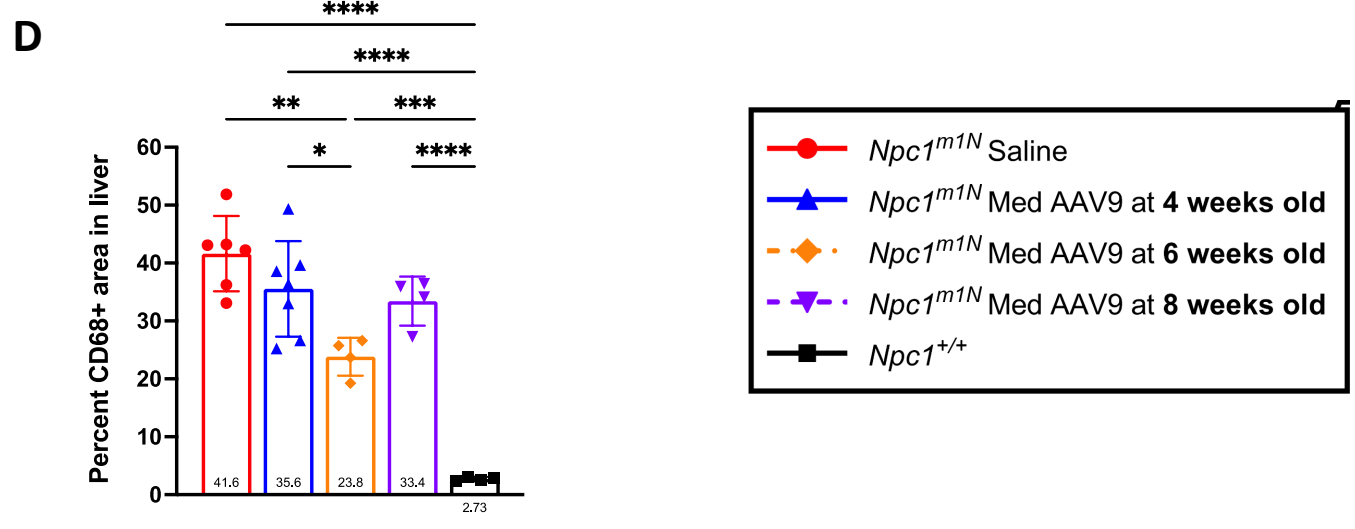
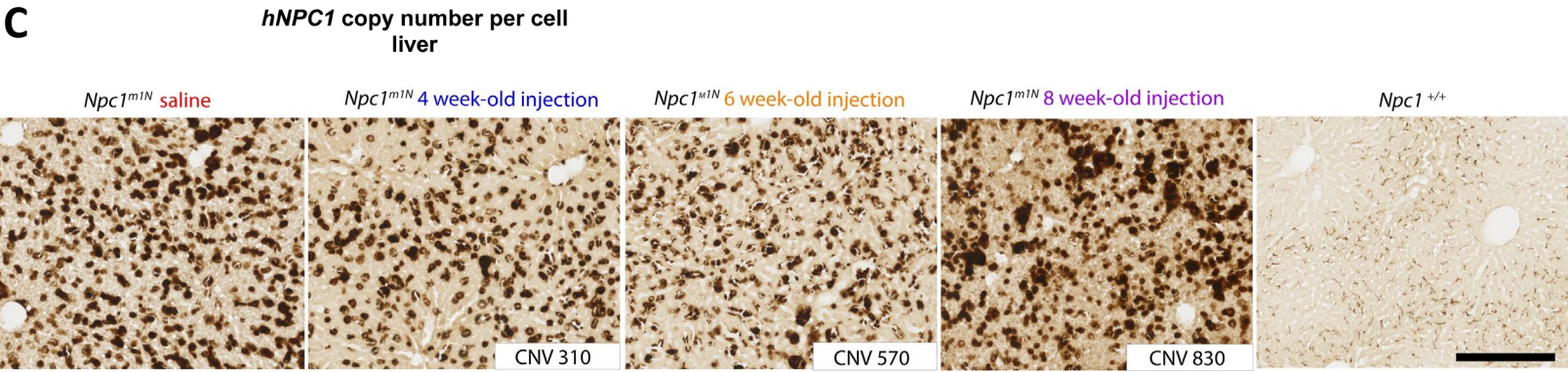
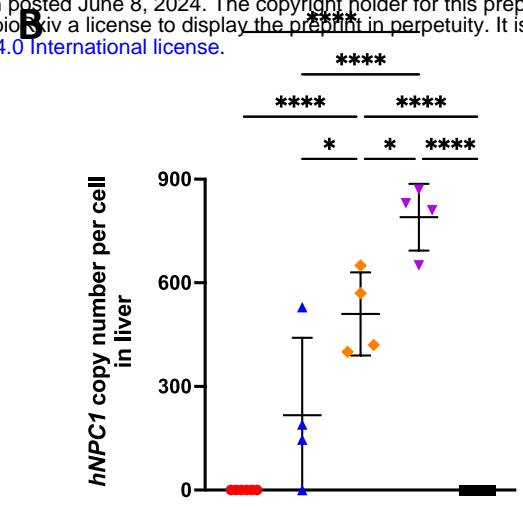
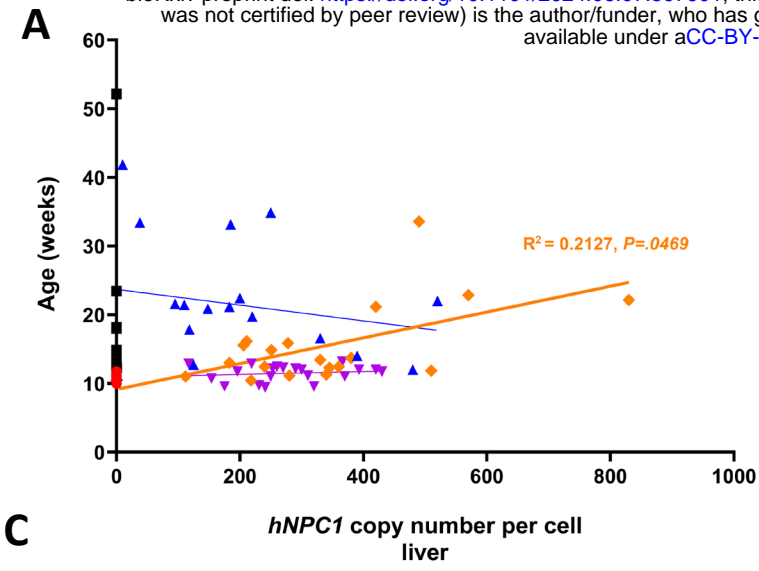


A

Treatment	Sample Size (n)	Significance (Tukey's Multiple Comparisons Test, 6-9 weeks)	Significance (Tukey's Multiple Comparisons Test, 9-12 weeks)
<i>Npc1<sup>m1N</sup></i> Saline	14		
<i>Npc1<sup>m1N</sup></i> Low	10	vs. Saline, P=0.9466	vs. Saline, P=0.9976
<i>Npc1<sup>m1N</sup></i> Medium	13	vs. Saline, P=0.0196 vs. Low, P=0.0069	vs. Saline, P<0.0001 vs. Low, P<0.0001
<i>Npc1<sup>m1N</sup></i> High	8	vs. Saline, P<0.0001 vs. Low, P<0.0001 vs. Medium, P = 0.0129	vs. Saline, P<0.0001 vs. Low, P<0.0001 vs. Medium, P = 0.0009
<i>Npc1<sup>+/+</sup></i>	21	vs. Saline, P<0.0001 vs. Low, P<0.0001 vs. Medium, P = 0.0003 vs. High, P = 0.9104	vs. Saline, P<0.0001 vs. Low, P<0.0001 vs. Medium, P<0.0001 vs. High, P = 0.5073

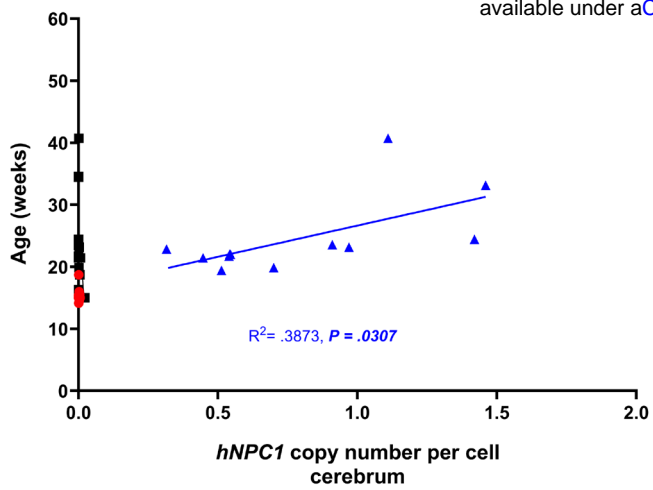
B

Treatment	Sample Size (n)	Significance (Tukey's Multiple Comparisons Test, 6-9 weeks)	Significance (Tukey's Multiple Comparisons Test, 9-12 weeks)
<i>Npc1<sup>m1N</sup></i> Saline	14		
<i>Npc1<sup>m1N</sup></i> Med AAV9 at 4 weeks old	13	vs. Saline, P<0.0001	vs. Saline, P<0.0001
<i>Npc1<sup>m1N</sup></i> Med AAV9 at 6 weeks old	20	vs. Saline, P=0.7788 vs. 4 weeks old, P=0.0010	vs. Saline, P=0.1909 vs. 4 weeks old, P=0.0002
<i>Npc1<sup>m1N</sup></i> Med AAV9 at 8 weeks old	20	vs. Saline, P=0.9984 vs. 4 weeks old, P<0.0001 vs. 6 weeks old, P=0.5116	vs. Saline, P>0.9999 vs. 4 weeks old, P<0.0001 vs. 6 weeks old, P=0.0747
<i>Npc1<sup>+/+</sup></i>	21	vs. Saline, P<0.0001 vs. Weaning, P=0.2389 vs. 6 weeks old, P<0.0001 vs. 8 weeks old, P<0.0001	vs. Saline, P<0.0001 vs. Weaning, P<0.0001 vs. 6 weeks old, P<0.0001 vs. 8 weeks old, P<0.0001

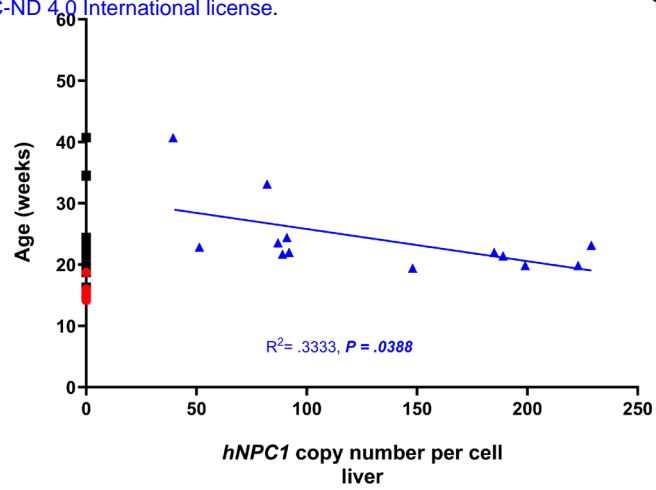




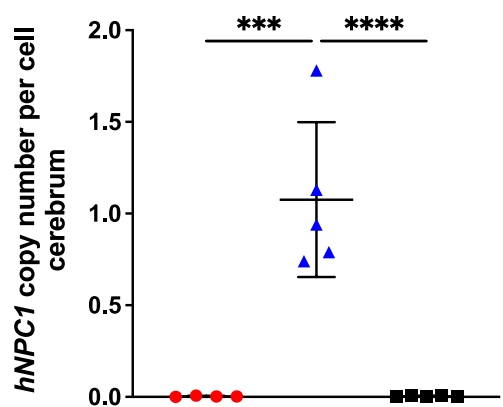
A



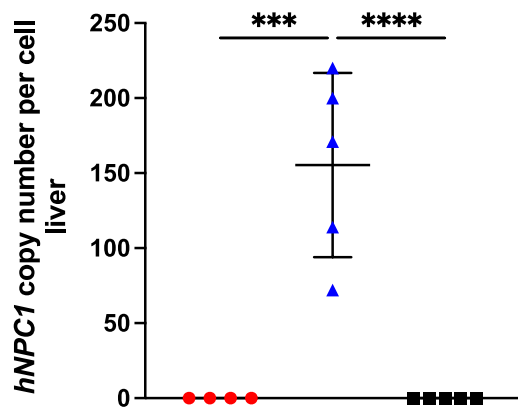
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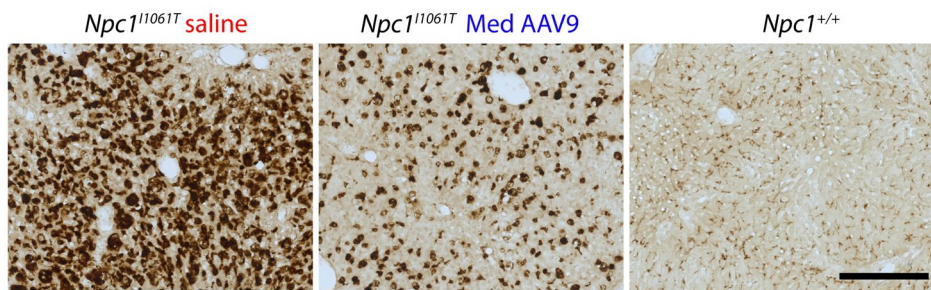
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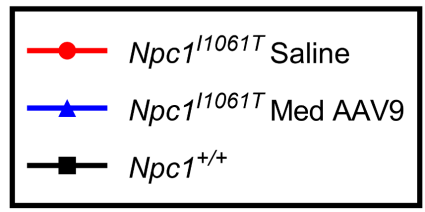
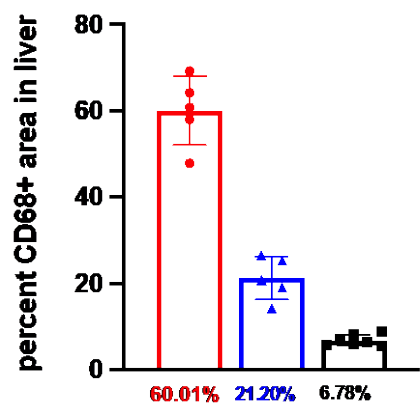
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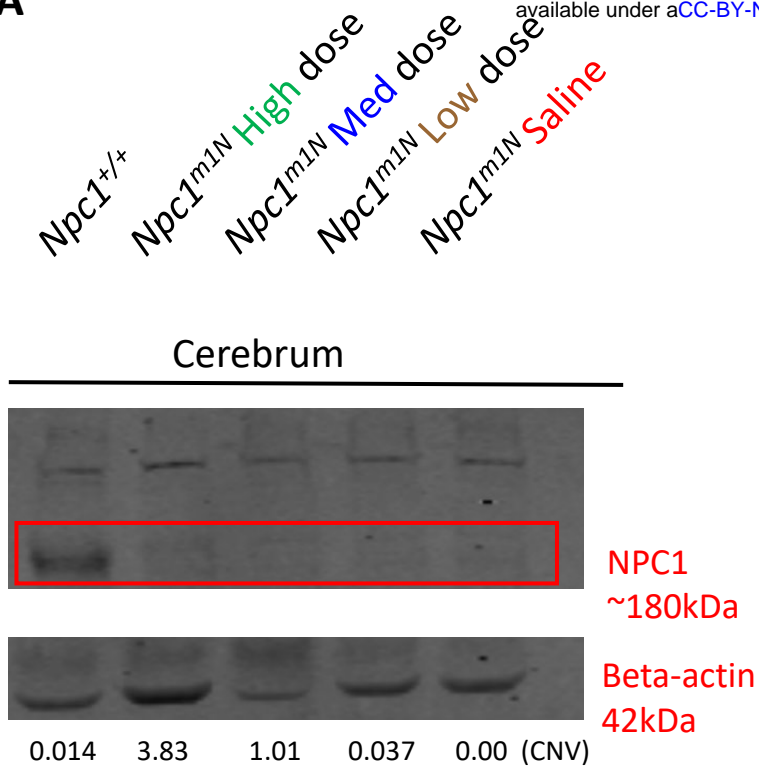
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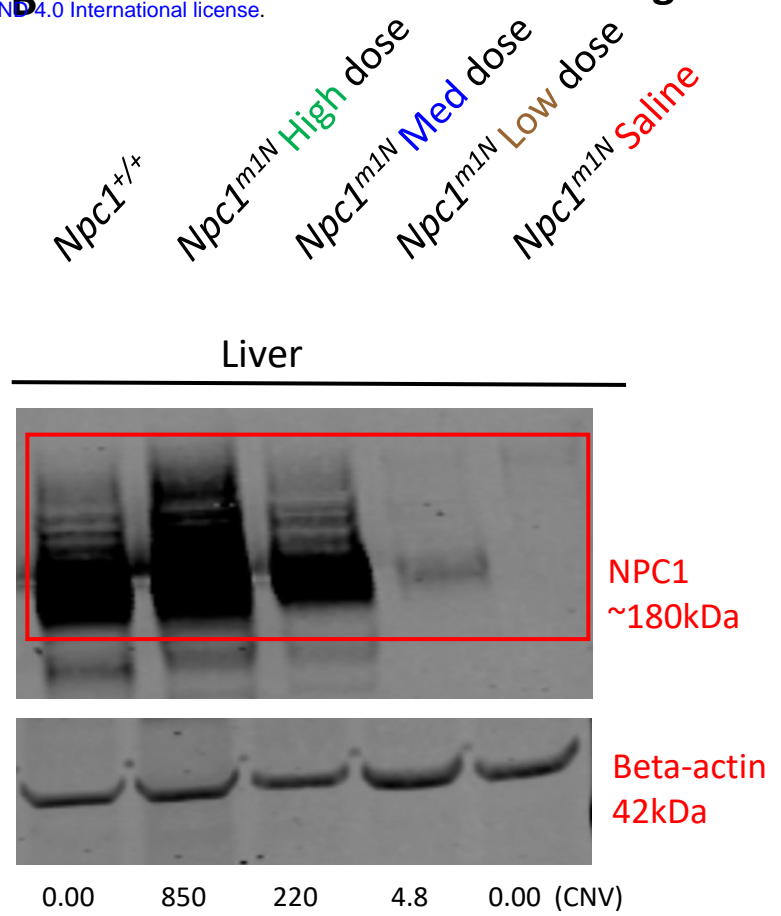
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A



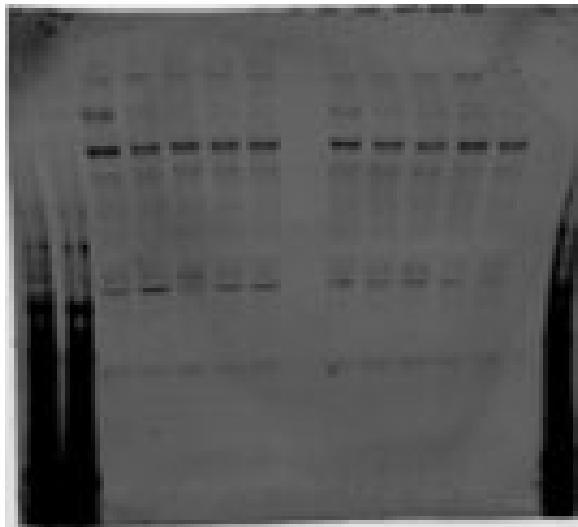
B



**Figure S6.** Representative blot for NPC1 quantification in cerebrum (Fig 2Aiii) and liver (Fig 2Biii). *hNPC1* copy number variation listed under blots for each sample.

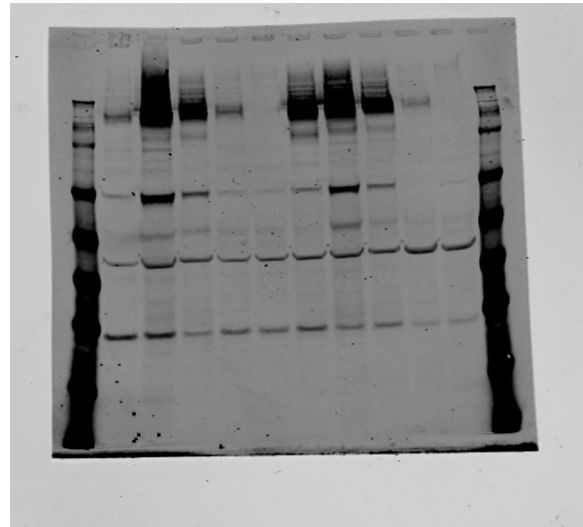
C

Original source data, cerebrum (NPC1, beta-actin; left half of blot enlarged).

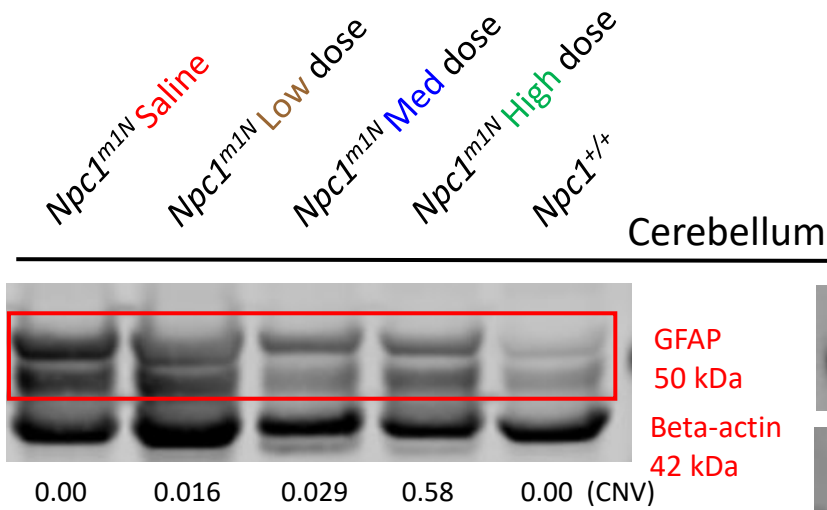


D

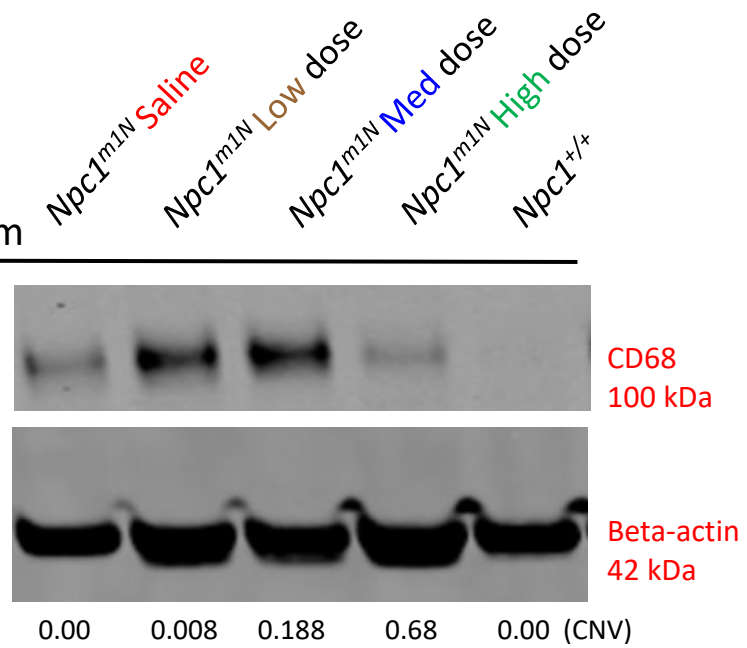
Original source data, liver (NPC1, beta-actin; right half of blot enlarged).



A



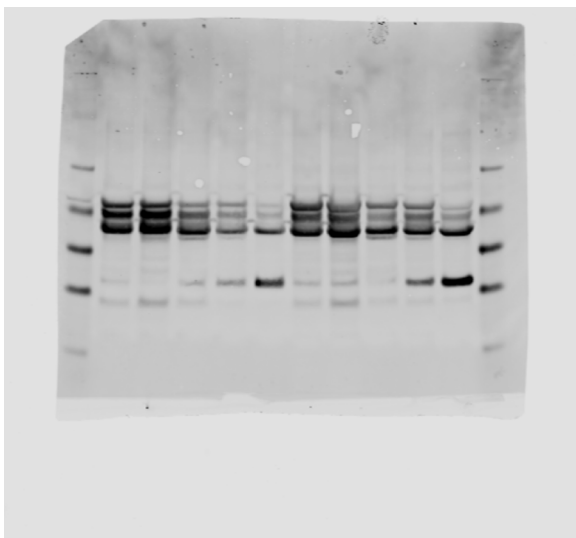
B



**Figure S7.** Representative blot for GFAP quantification (Fig 3C) and CD68 quantification (Fig 3E) in the cerebellum. *hNPC1* copy number variation listed under blot for each sample.

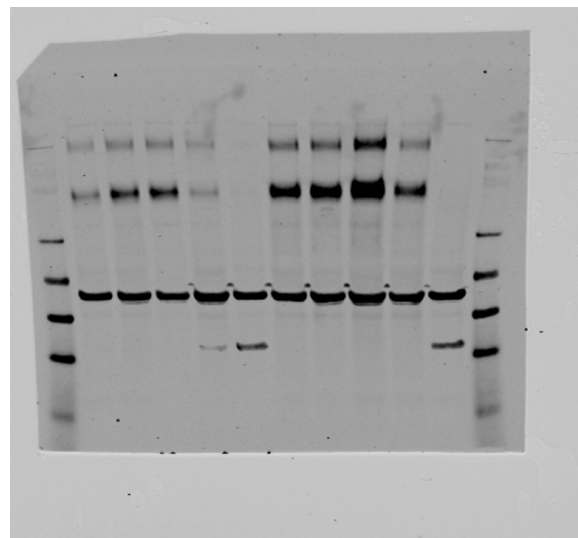
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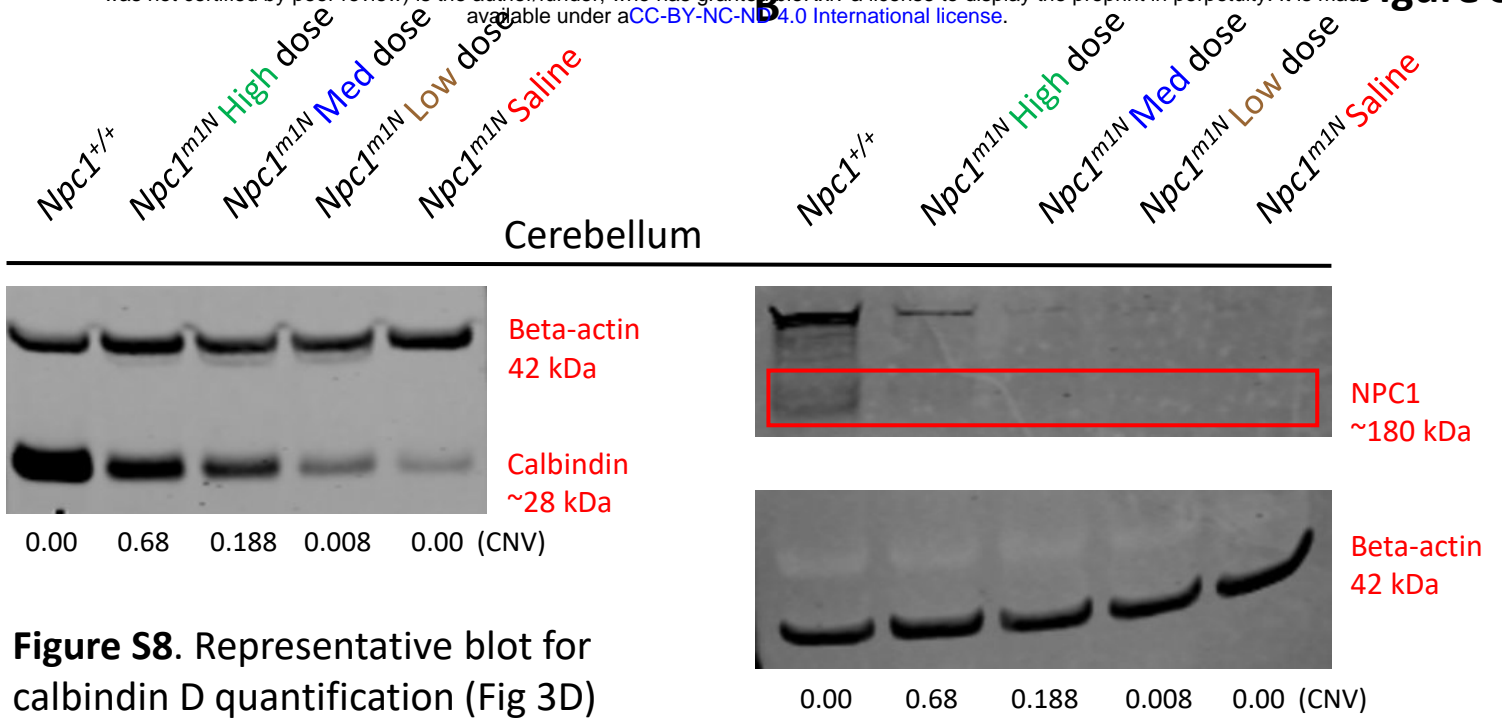
Original source data (GFAP, beta-actin; right half of blot enlarged).



D

Original source data (CD68, beta-actin; left half of blot enlarged).





**Figure S8.** Representative blot for calbindin D quantification (Fig 3D) and NPC1 quantification (Fig 3G) in the cerebellum. *hNPC1* copy number variation listed under blot for each sample.

**C** Original source data (Calbindin, beta-actin on right, NPC1, beta-actin on left).

