Supplemental Information

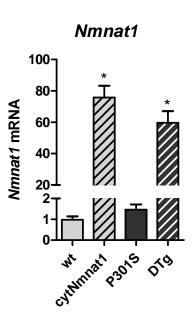


Fig. S1: cytNmnat expression is not significantly impacted by P301S transgene expression. Nmnat1 mRNA was quantified by qPCR in cortex samples. *p<0.05 by 1-way ANOVA with Bonferroni's post-test as compared to wt.

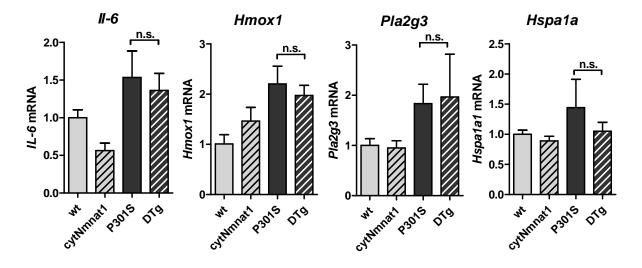


Fig. S2: Inflammation and oxidative stress response in P301S tau mice are not altered by cytNmnat1 overexpression. Cortex samples from wt, cytNmnat1, P301S, and DTg mice were subjected to qPCR analysis of mRNA expression of *II6* (inflammatory cytokine), *Hmox1* and *Pla2g3* (oxidative stress responsive genes) and *Hspa1a* (a subunit of the chaperone hsp70). A trends toward increase in all markers was observed in tau-expressing mice. However, no significant difference between P301S and DTg mice was observed for any of these markers. **p<0.05 by 1-way ANOVA as compared to wt. n.s. non-significant.