Characterization of Striatal Neuronal Loss and Atrophy in the R6/2 Mouse Model of Huntington’s Disease

January 6, 2014 · HD Models
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Abstract
Striatal neuronal degeneration and loss is an important feature of human Huntington’s disease (HD). R6/2 HD mice recapitulate many features of human HD including striatal atrophy. While striatal neuronal atrophy and loss is reported in R6/2 HD mice the degree of neuronal loss and the characteristics of cell body atrophy are unclear. We used stereological approaches to estimate whole striatal neuronal numbers and characterize changes in striatal neuronal size distribution. R6/2 HD mice had 126000 fewer neurons per striatum (~12% decline) at 12 weeks of age than wild-type litter-mates; differences were not present at 5 weeks. Analysis of striatal neuronal numbers per cell body size category revealed declines in neuron numbers in the size ranges 550-1050 µm³ suggesting that larger striatal neurons are more susceptible to atrophy or loss in late stages of disease. R6/2 HD mice have a striatal neuronal loss phenotype. As striatal neuronal loss in human HD is dramatic, neuronal loss in R6/2 striatum provides an important late-stage outcome measure for study of disease modifying interventions.

Funding Statement
This research was funded in part by NIH grants 5P20RR015640-10 and NIH R01 NS079450-01. Lindsay Dodds was supported by an INBRE undergraduate student summer fellowship (NIH P20RR016474 and NIH P20GM103432).

Introduction
Human Huntington’s disease (HD) is characterized by slowly progressive brain atrophy primarily affecting striatum and cerebral cortex that develops in the pre-clinical phase of the disease process. There is considerable evidence that atrophy of striatum results from a combination of neuronal degeneration and loss. Vonsattel et al reported loss of 50% of caudate neurons in neuropathology grade I and 95% in grade IV human disease. Neurons within the striatum comprise ~95% medium spiny neurons which degenerate early in HD and remaining interneurons which are less susceptible to degeneration.

Numerous genetic mouse models of HD exist and these have proven invaluable for studying disease mechanisms and therapies. R6/2 mice express exon 1 of the human huntingtin gene containing an expanded CAG repeat. These mice experience rapid disease progression with symptoms beginning around 6 weeks of age, large behavioral declines by 12 weeks, and death as early as 10 weeks of age in some colonies. R6/2 mice recapitulate many features of human HD including progressive brain atrophy involving neostriatum and cerebral cortex, weight loss, motor dysfunction and accumulation of mutant huntingtin aggregates in brain neurons. While these mice express only a fragment of human mutant huntingtin evidence has been provided that this fragment occurs endogenously in mouse HD.

YAC128 HD mice express full-length human mutant huntingtin from a yeast artificial chromosome. They have a slowly progressive HD-like disease process. Slow and colleagues reported a 15% decrease in striatal volume at 9 months of age, and a 15-18% reduction in striatal neuronal estimates at 12 months of age in these mice. Knock-in models of HD also exhibit late-onset neurodegeneration. For example, in the CAG140 mouse HD model 38% and 40% reductions in striatal volume and striatal neuron estimates are reported, respectively, in 20-26-month-old animals. While fragment mouse models of HD (e.g. R6/2) are genetically less accurate than full-length models their rapid disease progression and HD-like phenotype has resulted in their continued use.

Several approaches have been used to demonstrate striatal neuronal degeneration in HD models. R6/2 HD mice have increased activity of neuronal caspase 1 and 3 and changes in Bcl-2 family members supportive of activation of striatal neuronal apoptosis. One study reported significant mitochondrial abnormalities in neurons of R6/2 mice. Findings from TUNEL labeling for DNA nicks have however been variable. Several studies have demonstrated the presence of neuronal cell body atrophy in the striatum of R6/2 HD mice. However, we are aware of only one study that has estimated striatal neuronal numbers in R6/2 mice as a means to quantify neurodegeneration. Stack et al estimated neuronal numbers in the anterior 2/3 of the striatum, to the level of the anterior commissure, and found significantly fewer neurons at 12-weeks of age in...
R6/2 mice compared to wild-type littermates. The purpose of the current study was to complete a detailed stereologic analysis of the entire striatum, in particular characterizing neuronal numbers and size changes with disease progression.

Methods

All procedures were pre-approved by the University of Wyoming Animal Care and Use Committee (IACUC). R6/2 HD mice were bred and maintained as previously described\textsuperscript{5}. CAG repeat sizes were determined from genomic DNA by Laragen Inc. (Culver City, CA) and averaged 181 with a standard deviation of 6. Mice were sacrificed by deep anesthesia then intra-cardiac perfusion with 4% paraformaldehyde and processed for stereology as described\textsuperscript{15}. In brief, brains were sectioned coronally to include the entire striatum at 40 \( \mu \)m thickness; every 12\textsuperscript{th} section was mounted and stained using the thionin method. We used the unbiased optical dissector method to estimate striatal neuronal numbers\textsuperscript{16}. We used a counting frame size of 40 x 40 \( \mu \)m and a grid size of 500 x 500 \( \mu \)m. During the optical dissector procedure we also estimated striatal neuronal cell body volume using the 2D nucleator with five isotropic uniform random sections per cell. All slides were coded so the operator was unaware of the genotype. The numbers of mice per group were: wild-type 5-weeks n=15; HD 5-weeks n=13; wild-type 12 weeks n=9; and HD 12 weeks n=10. The average number of neurons / mouse analyzed was 384 for 5-week mice (range: 245-521) and 359 for 12-week mice (range: 272-436). For each mouse, neurons were binned into 50 \( \mu \)m\textsuperscript{2} cell body size categories using Microsoft excel software. The estimated total number of striatal neurons in each size category in wild-type and HD mice was then analyzed using the students t-test in SAS software version 9.2 (Cary, NC, USA).

Results

R6/2 HD mice had 12.7% fewer striatal neurons at 12-weeks of age (\( p=0.035 \)) (Fig. 1A). Wild-type mice had a mean of 993891 ± 38483 (SEM) neurons; R6/2 HD mice had a mean of 868043 ± 38483 (SEM). To determine if this difference was due to neuronal loss or developmental effects we studied 5-week-old mice. As shown in Fig. 1B there were no differences at 5-weeks of age indicating the presence of neuronal loss in R6/2 mouse striatum occurring between 5 and 12 weeks. Use of the nucleator stereologic tool has been reported in numerous studies to characterize changes in striatal neuronal cell body volume in HD mice; however, distribution of neuronal sizes and number estimates by cell size were not determined (e.g. [12,13]). We analyzed neuronal population estimates on a cell body size basis (Fig. 2). At 12 weeks, the most significant finding was a loss of neurons in the size range 550-1050 \( \mu \)m\textsuperscript{3} (Fig. 2A). Within smaller neuron size categories (200-500 \( \mu \)m\textsuperscript{2}) neuronal population mean estimates were increased in R6/2 mice but this did not approach significance (Fig. 2A). At 5 weeks, there were no significant differences in neuron number estimates per size category (Fig. 2B).

While there is abundant biochemical and morphologic evidence to support neuronal degeneration in R6/2 mouse striatum\textsuperscript{10,11,17}, we are not aware of reports of morphologic evidence of late-stage neuronal death events in R6/2 striatal neurons. Our findings, and those of others, suggest that while the process of neuronal degeneration is gradual, neuronal death and removal is rapid. We estimated the number of striatal neurons in the process of death when sacrificed at 12-weeks of age. To do this we assumed that neuron death in R6/2 mice begins at 60 days of age\textsuperscript{14} and that the process is linear across time (a constant number of neurons dying / day). Based on this, we calculated that R6/2 HD mice lose ~6909 neurons / day / striatum. In one rodent model of acute neuronal injury cell death was detected ~6 hours post-insult\textsuperscript{18}. In one study phagocytosis of apoptotic neurons occurred within 1.5 hours of introduction of microglia into cultures\textsuperscript{19}. Based on these findings we assumed that in R6/2 mice the process of morphologically detectable neuronal death and removal lasts 8 hours. From this we calculated that a 20 \( \mu \)m section of striatum taken at the level of the anterior commissure of a 12 week R6/2 mouse would contain ~4 dying neurons. Therefore, based on our estimates, the kinetics of neuronal degeneration, loss and removal underlies the difficulty of identifying neuronal death morphologically. Stereologic approaches to quantifying neurons, as used here, can overcome these difficulties.
Striatal neuronal loss occurs in R6/2 HD mice. A. R6/2 HD mice have 12.7% fewer neurons than wild-type litter-mate mice at 12 weeks of age. p=0.035, B. R6/2 HD mice do no differ from wild-type litter-mate mice in striatal neuronal number estimates at 5 weeks of age. P-value: *=p<0.05

**Fig. 1: Striatal neuronal loss occurs in R6/2 HD mice**

Striatal neuronal atrophy and loss in R6/2 HD mice. A. R6/2 HD mice have fewer neurons in the cell body size range 550-1050 µm³ at 12 weeks of age. B. There are no significant differences in neuronal number estimates per cell body size category in 5-week-old mice. Round circles = R6/2 HD; closed circles=wild-type. P-values: *=p<0.05, **=p<0.01

**Fig. 2: Striatal neuronal atrophy and loss in R6/2 HD mice.**

Numerous studies have documented a reduction in striatal neuronal cell body size in R6/2 HD mice. Here we took a new approach to analysis by determining estimates of neuronal numbers in specific size categories. Our analysis cannot clearly reveal in what population neuronal loss occurs in R6/2 striatum as decreased numbers of neurons in the size range 550-1050 µm³ (Fig. 2A) could result from atrophy, loss or a combination. However, the results do suggest that larger medium spiny neurons are more vulnerable to degeneration in R6/2 HD.

Taken together, our findings corroborate and extend those reported previously. Our analysis of striatal neuron number within whole striatum reveals a 12% loss of neurons in R6/2 mice which is less than the 26% loss reported in an analysis based on examination of the anterior striatum to the level of the anterior commissure, which is about 2/3 of the striatum. This discrepancy could be for a number of reasons that includes differences in neuronal loss in whole striatum versus anterior striatum and differences in the respective R6/2 mouse models including CAG size and background genetic effects due to the backcrossing approach to colony maintenance. Importantly, however, all mouse models of HD studied so far have significantly less striatal neuronal loss than the 90% reported in advanced human HD. While striatal neuronal loss in R6/2 mice and other mouse HD models is modest, the presence of dramatic loss in human HD brain suggests that estimation of striatum neuronal number should be an important outcome in HD mice which can be used to evaluate therapies in pre-clinical studies. Finally, we undertook a sample size analysis to determine the number of mice required to detect rescue of neuronal loss in R6/2 mice in a hypothetical treatment study. Parameters used were: 80% power, a significance level of 0.05, standard deviations proportional to the mean and a one-tailed t-test. Based on these parameters we calculated that for 50% and 100% rescue of striatal neuronal loss it would be necessary to have group sizes of 10 and 5, respectively. While stereologic approaches to estimate...
neuronal loss can be labor intensive, our calculations indicate that only modest group sizes are required. Analysis of neuronal numbers may be especially valuable for interventions that are protective in mice and when further characterization of protective effects is needed prior to considering human testing.

Acknowledgements

Lindsay Dodds and Jianfang Chen contributed equally to this work. We thank Dr. Kenneth Gerow, Department of Statistics, University of Wyoming, for assistance with the sample size analysis.

References

