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Association of CAG Repeats With Long-term Progression in Huntington Disease

Langbehn, D. R., Stout, J. C., Gregory, S., Mills, J.A., Durr, A., Leavitt, B. R., Roos, R. A. C.,
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Landwehrmeyer, B., Scahill, R. I., & Tabrizi, S. J.

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MJC Presentation

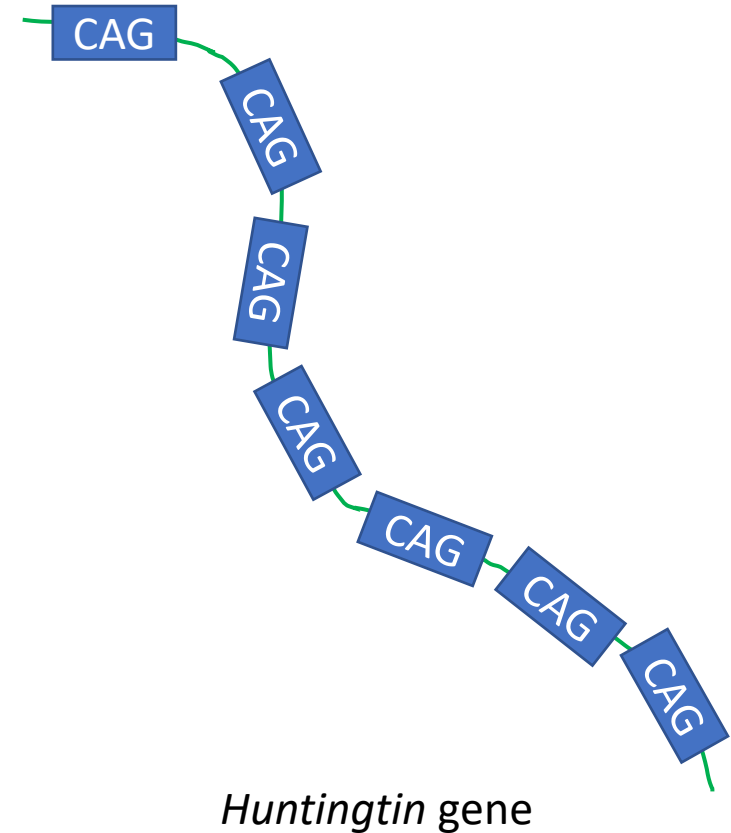
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Abstract Abstracted

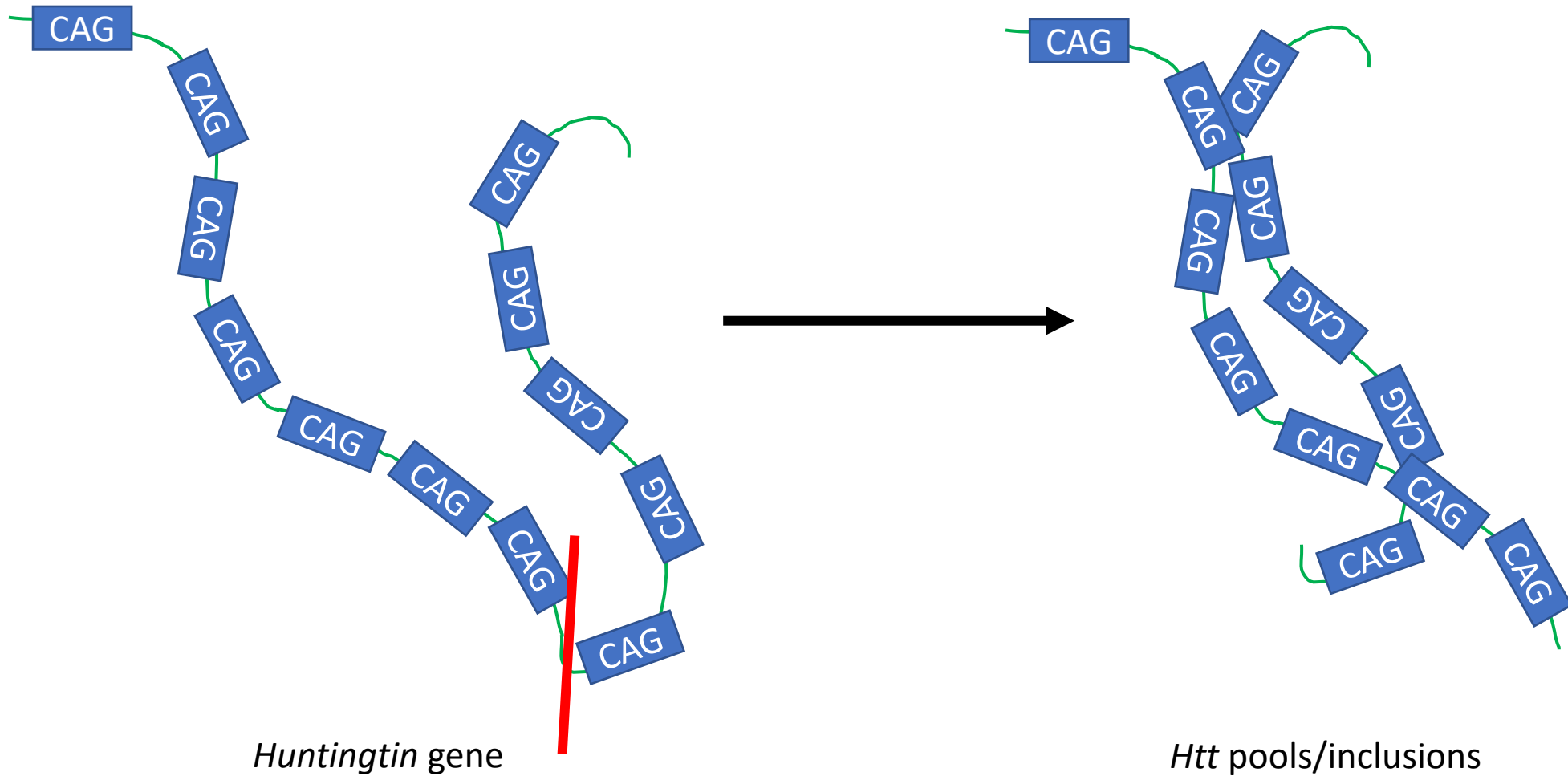
- Huntington's disease is neurodegenerative disease caused by a single mutation in the Huntingtin gene
- The number of CAG repeats can be used to predict the severity of degeneration in the brain as well the age of disease onset
- Disease progression can be defined in terms of a number of variables (including performance on tests and brain volume), but the most basic one is time
- Langbehn et al. found that putamen and caudate volumes best tracked disease progression at any given time

Introduction

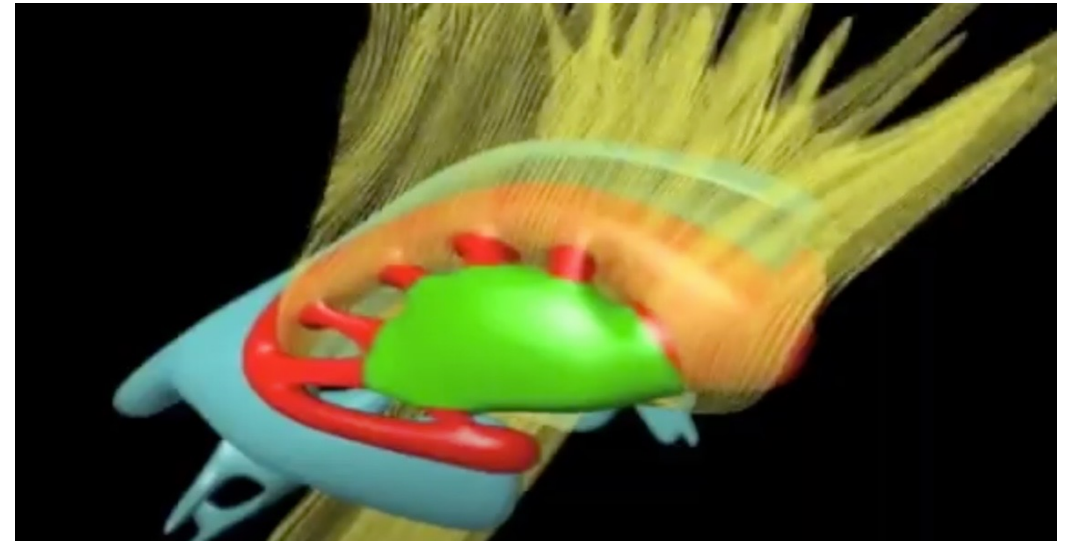
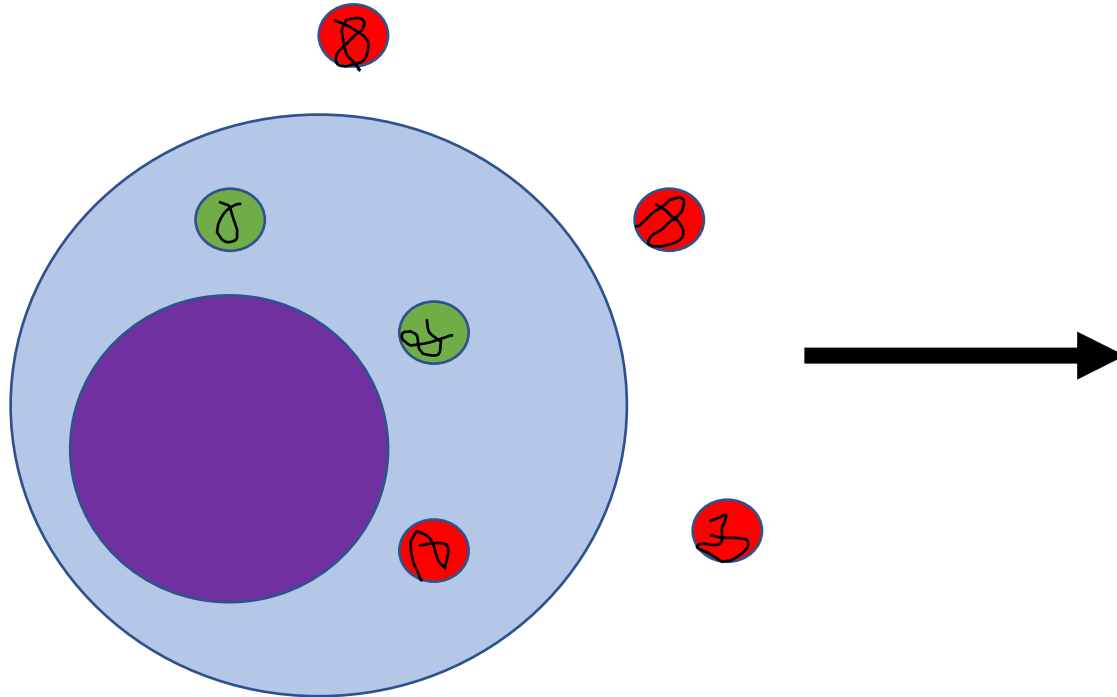
- Huntington's disease is a neurodegenerative disease characterized by expansion of the CAG trinucleotide sequence in the *huntingtin* gene of chromosome 4
- Therefore, Huntington's disease is inherently a genetic disease



Introduction



Introduction



*Intracellular and extracellular inclusions develop
(Jimenez-Sanchez et al., 2017)*

*↓ brain volume in the striatum
(Alila Medical, 2021)*

Introduction

- The primary clinical markers of the disease are cognitive and motor deficits
 - Perform worse on a tapping task
 - Perform worse on a Stroop word reading task
- The expression of the disease symptomatically based on the genetic underpinning is known as *penetrance*
 - 10-26 repeats → no penetrance
 - 26-25 repeats → intermediate penetrance
 - 39-39 repeats → reduced penetrance
 - 40+ repeats → full penetrance

| | | | | | |
|--------|--------|--------|--------|--------|--------|
| Red | Red | Blue | Green | Red | Red |
| Yellow | Green | Green | Green | Yellow | Green |
| Blue | Blue | Yellow | Yellow | Blue | Blue |
| Green | Red | Red | Blue | Green | Red |
| Green | Blue | Green | Yellow | Green | Green |
| Yellow | Green | Blue | Blue | Blue | Yellow |
| Blue | Yellow | Red | Green | Red | Blue |

Introduction

- Langbehn et al. claim that few disease characteristics can be determined using just the CAG repeat length and age
- The goal of the research was to determine the clinical and morphometric implications of CAG repeat length
- Hopefully, the results would help determine:
 - 1. Which variables are best associated with CAG repeat length and age (highest R^2)
 - 2. How the progression of Huntington's disease looks like (Quadratic? Linear? Flat?)

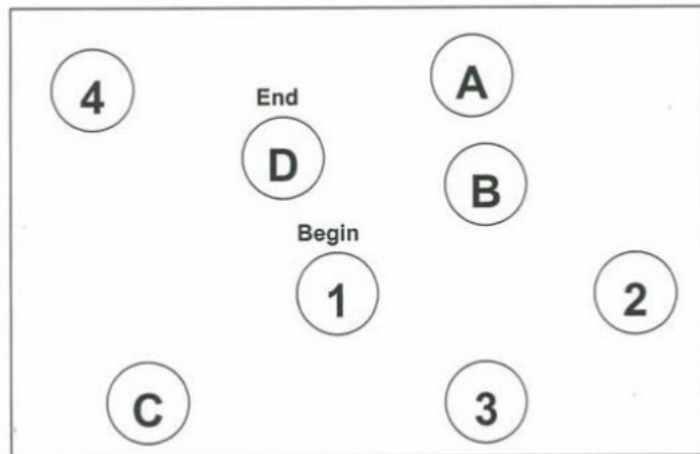
Methods

- The researchers did not sample individuals themselves
- Rather, they obtained data through the TRACK-HD/TRACK-ON studies

| TRACK-HD | TRACK-ON |
|--|---|
| <ul style="list-style-type: none">• four academic institutions first recruited 123 healthy controls, 120 pre-HD carriers, and 123 with early HD subjects | <ul style="list-style-type: none">• a follow up of TRACK• still occurring• maximum time of consideration: 6 years |
| Combined | |
| <ul style="list-style-type: none">• 2065 patient visits• 443 participants• Used UHDRS variables + morphometric data | |

Methods

- UHDRS – Clinical
 - A clinical battery of tests used to assess disease severity
 - Involves motor, cognitive, and neuropsychiatric tasks/tests
 - 74 total items



- Morphometric data (MRI)
 - Brain volumes (caudate and putamen, total brain volume)
 - Given that people naturally have different-sized heads, the **ratios** of these volumes to the total intracranial volume

$$\text{Volume Considered} = \frac{\text{Actual Volume}}{\text{Total Intracranial Volume}}$$

Methods

- Principal component analysis was used to derive summary scores for clinical and morphometric data
 - *“the weighted sum of the original set of variables that has the highest possible mean correlation with those same variables”*
 - That statistics themselves are described using linear algebra

$$\mathbf{w}_{(1)} = \arg \max_{\|\mathbf{w}\|=1} \left\{ \sum_i (t_1)_{(i)}^2 \right\} = \arg \max_{\|\mathbf{w}\|=1} \left\{ \sum_i (\mathbf{x}_{(i)} \cdot \mathbf{w})^2 \right\}$$

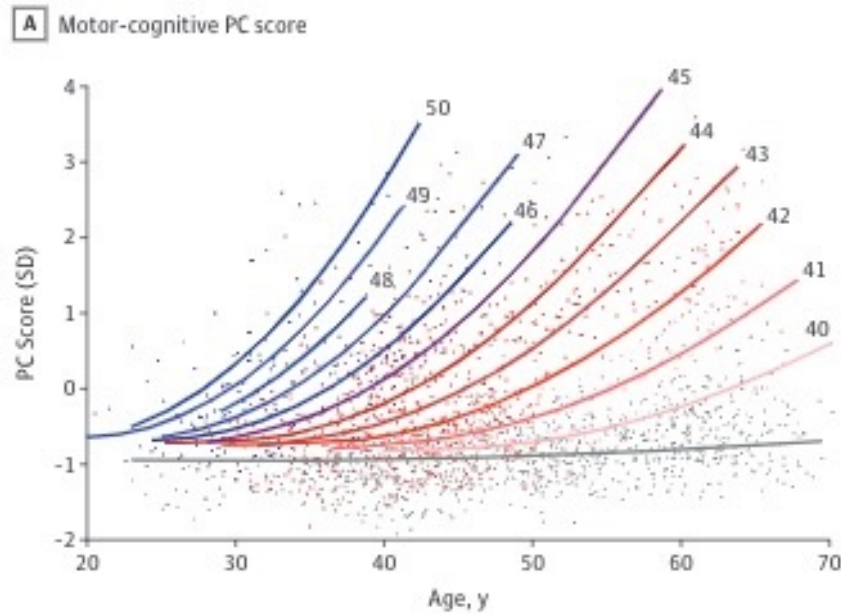
$$\mathbf{w}_{(1)} = \arg \max_{\|\mathbf{w}\|=1} \left\{ \|\mathbf{X}\mathbf{w}\|^2 \right\} = \arg \max_{\|\mathbf{w}\|=1} \left\{ \mathbf{w}^T \mathbf{X}^T \mathbf{X} \mathbf{w} \right\}$$

$$\mathbf{w}_{(1)} = \arg \max \left\{ \frac{\mathbf{w}^T \mathbf{X}^T \mathbf{X} \mathbf{w}}{\mathbf{w}^T \mathbf{w}} \right\}$$

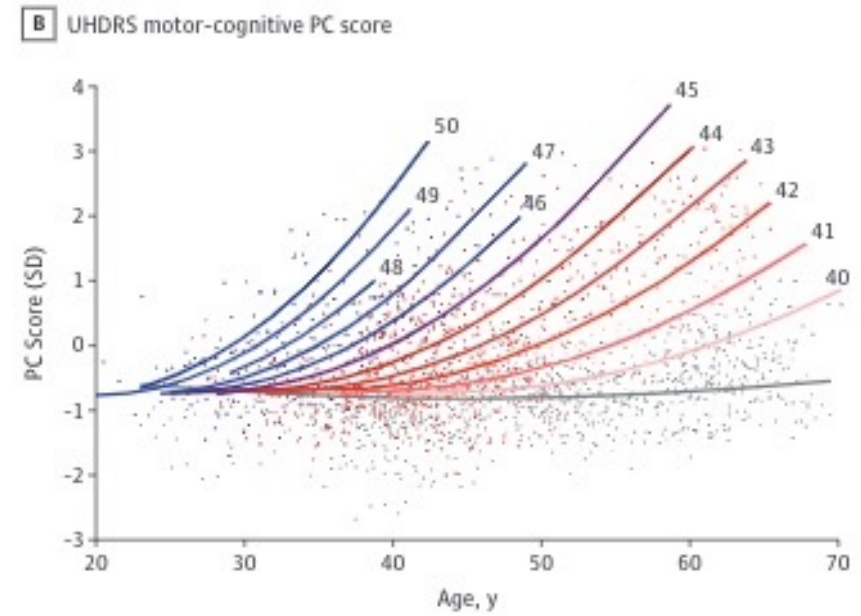
Results

| Variable | Motor-Cognitive Score | UHDRS Motor-Cognitive Score | White Matter-Ventricle Score | Caudate-Putamen Score |
|---|-----------------------|-----------------------------|------------------------------|-----------------------|
| Symbol Digit Modalities Test, No. correct | -0.795 | -0.903 | NA | NA |
| Stroop word reading, No. correct | -0.770 | -0.879 | NA | NA |
| Spot the Change 5 s, corrected for guessing | -0.582 | NA | NA | NA |
| UHDRS total motor score | 0.872 | 0.891 | NA | NA |
| Paced tapping 3-Hz SD of intertap intervals, log | 0.808 | NA | NA | NA |
| Q-Motor speeded tapping intertap interval, mean | 0.829 | NA | NA | NA |
| Q-Motor speeded tapping interonset interval, log SD | 0.885 | NA | NA | NA |
| Q-Motor speeded tapping tap duration, log SD | 0.889 | NA | NA | NA |
| Q-Motor speeded tapping interonset interval, log mean | 0.918 | NA | NA | NA |
| Putamen volume, ratio to ICV | NA | NA | NA | -0.947 |
| Caudate volume, ratio to ICV | NA | NA | NA | -0.947 |
| Total brain volume, ratio to ICV | NA | NA | -0.926 | NA |
| Ventricle volume, ratio to ICV | NA | NA | 0.818 | NA |
| White matter volume, ratio to ICV | NA | NA | -0.870 | NA |
| Gray matter volume, ratio to ICV ^b | NA | NA | NA | NA |
| Variance explained by the first PC, % | 67.6 | 79.4 | 76.1 | 89.7 |

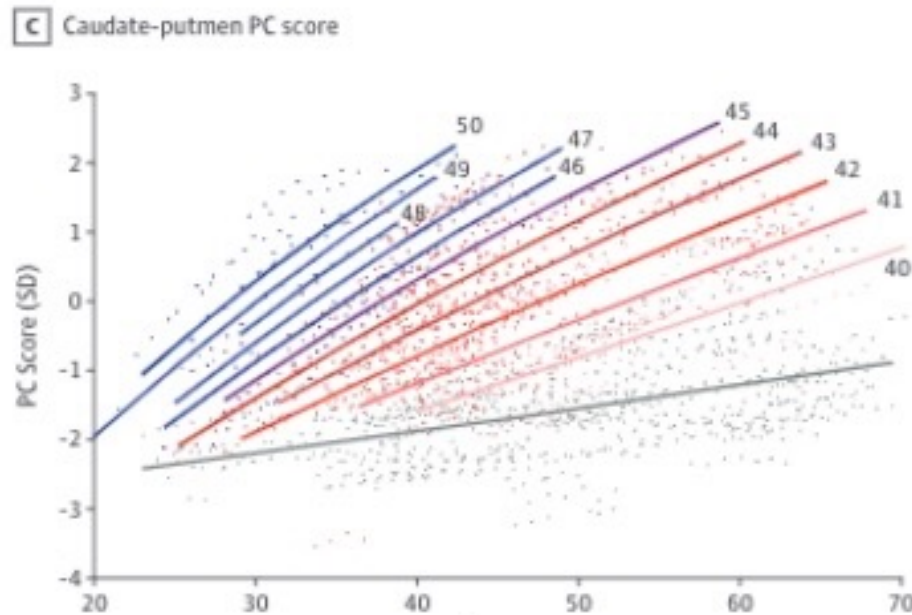
Results



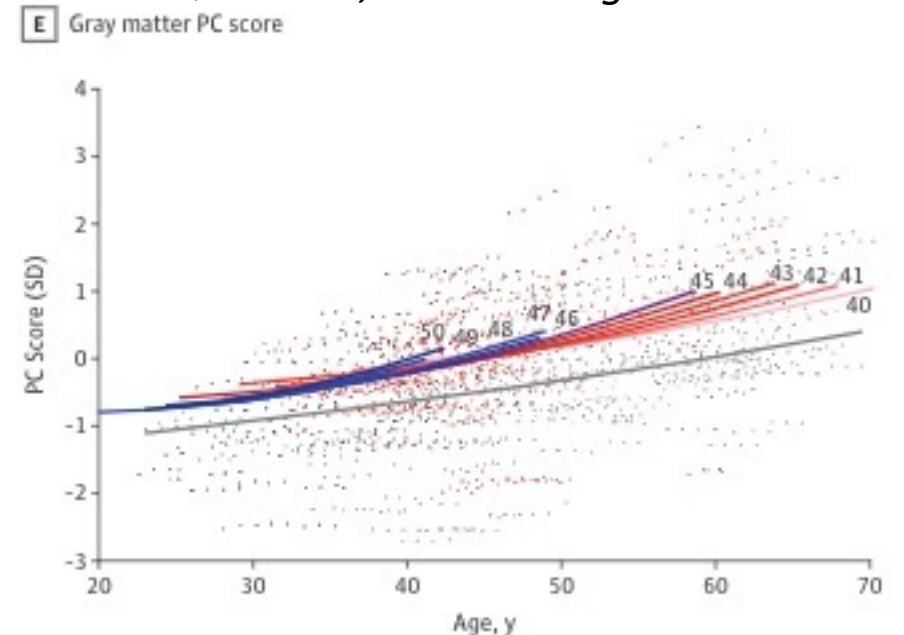
Quadratic, Accelerating



Quadratic, Accelerating

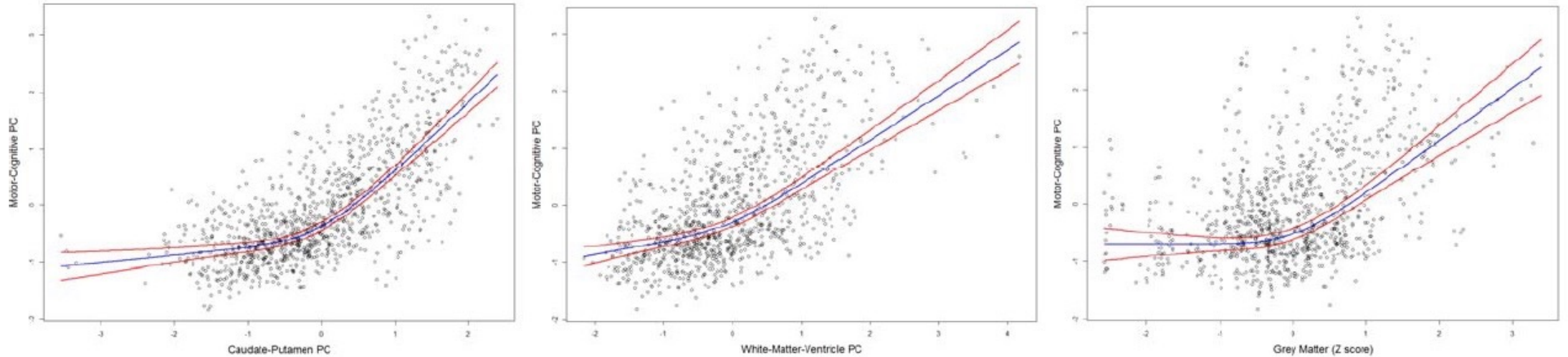


Quadratic, Accelerating



Non-linear, low slope

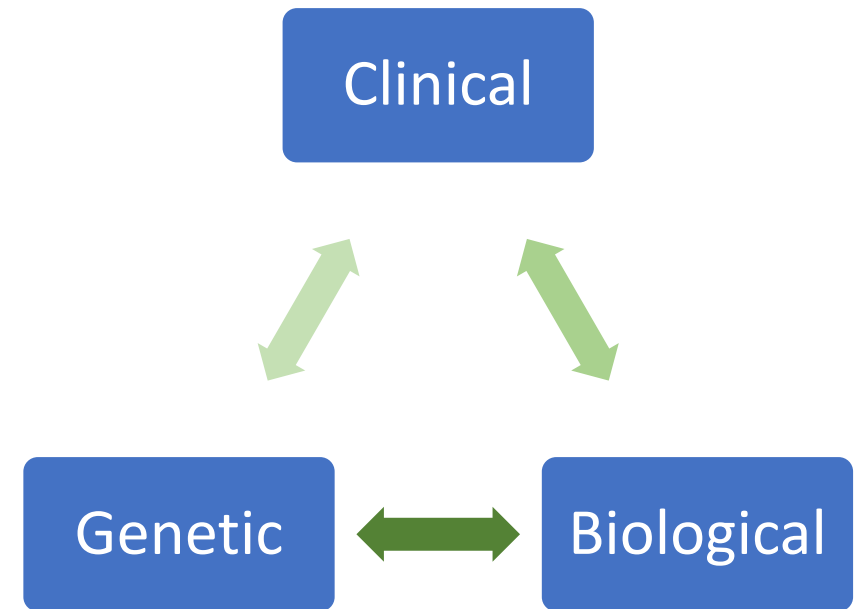
Results



- Previous graphs depicted motor-cognitive, caudate-putamen, and gray matter scores as a function of time
- These graphs indicate the relationship between the PCs themselves
- The inflection point at PC = 0 indicates shift from healthy to HD subjects

Conclusion

- Phenotype declines in a CAG-dependent manner
 - However, different acceleration rates between CAG repeat lengths indicate that a relationship is present
- Clinical predictors do not as strongly track disease progression as morphometric predictors do
- Variance can be decreased using a smaller subset of variables
- Neurodegeneration in the caudate and putamen is linear over time



Discussion Questions

- Why do you think brain matter volume tracked age in a quadratic manner?
- Why do you think neurodegenerative diseases have no cure? (Keep in mind genetics)
- What directions do think neuroscience/neurology should take in the future?

References

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