A digital motor score for sensitive detection of disease progression in early manifest Huntington's disease

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What does this mean for the Huntington's disease (HD) community?

The number of participants required for a clinical trial is directly related to how sensitively disease progression can be tracked. Furthermore, the larger a clinical trial, the longer it takes to recruit participants. The HD digital motor score has better sensitivity to change compared with standard clinical assessments, and may thus enable smaller/faster early-stage trials (sample size reduction by >75% possible). This means that, within the same timeframe, more therapies can be evaluated in a rare disease with a limited pool of potential study participants.

Conclusions

• A score built from four digital motor assessments showed a markedly increased sensitivity to change (STC) in people with Huntington's Disease Integrated Staging System (HD-ISS) Stage 3 and even HD-ISS Stage 2 (diagnostic confidence level 4); the STC at Week 20 was comparable to the STC of composite Unified Huntington's Disease Rating Scale at Week 69. Thanks to the large digital dataset collected across the Roche HD programme, development and validation of the HD digital motor score were performed on datasets from different clinical trials. Together with the good measurement properties observed, this increases confidence in the generalisability of the score.

Objective

To evaluate whether a combination of select features derived from motor assessments from the Roche Huntington's disease (HD) digital monitoring platform (DMP) into a single HD digital motor score (HDDMS) allows quantification of disease progression in people with Huntington's Disease Integrated Staging System (HD-ISS) Stage 2 and HD-ISS Stage 3 HD with higher sensitivity than standard clinical assessments.

Background

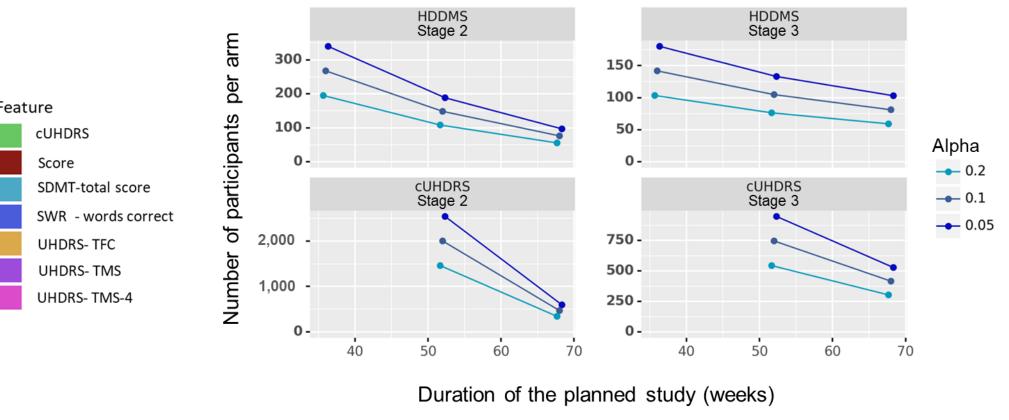
- The development of novel therapeutics for HD requires sensitive assessments of symptoms, especially for individuals with early-stage HD. The current standards, such as the composite Unified Huntington's Disease Rating Scale (cUHDRS) are valid, reliable and able to detect changes in large Phase III studies.¹
- However, in-clinic assessments are influenced by the interaction between rater and study participant, by their ordinal scale intervals, as well as day-to-day fluctuations of symptoms.
- Here, we propose an HDDMS, constructed •

Results

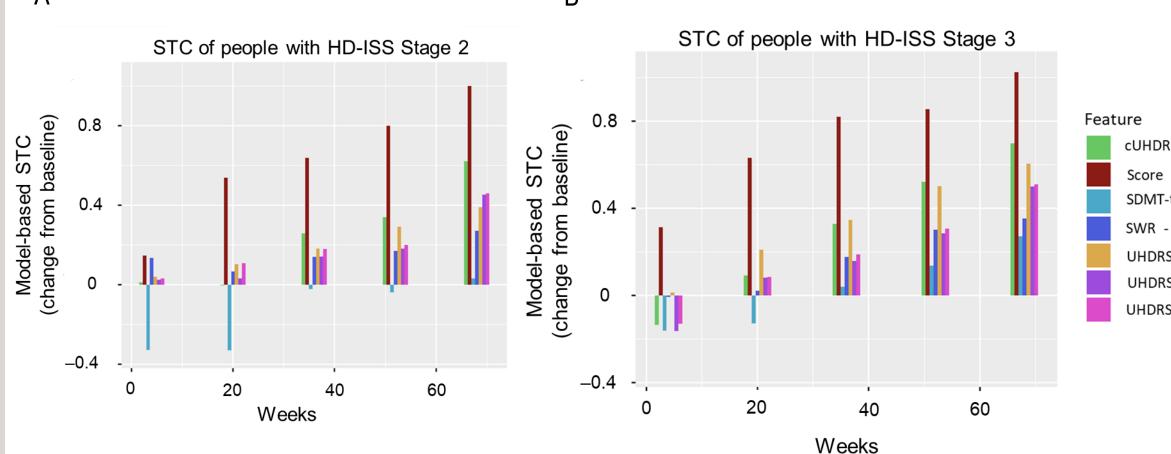
The HDDMS has a high sensitivity to change, resulting in lower sample sizes of people with HD-ISS Stage 2 or 3 in the GENERATION HD1 study.

Figure 1. STC of the HDDMS versus clinical assessments in people with HD-ISS Stage 2 or 3

Figure 2. Sample size calculation for cUHDRS and the HDDMS



Sample sizes for power=0.8 and 40% effect size. Sample size calculation for



from active digital motor assessments performed frequently at home, to sensitively assess progression of HD in both early and late disease stages.

Methods

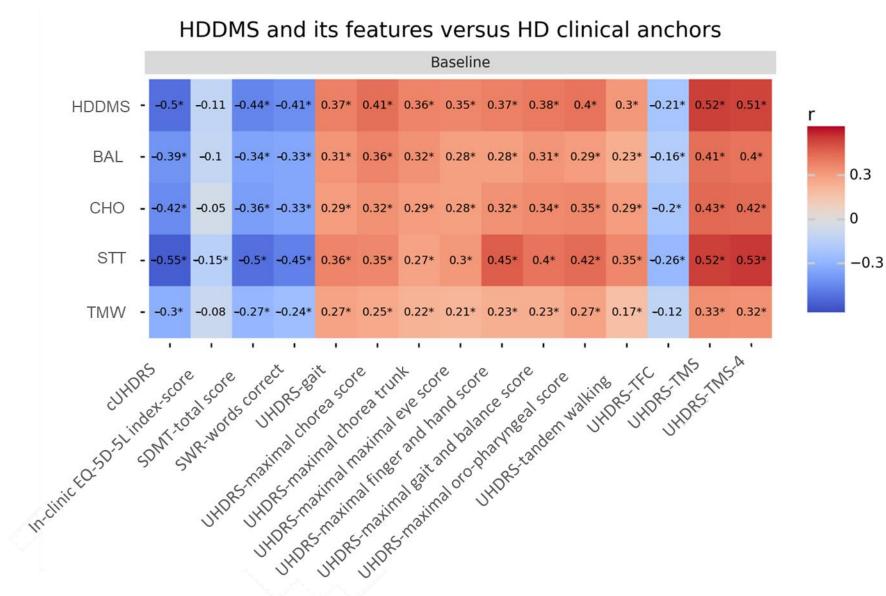
- We combined the digital features collected \bullet with the Roche HD DMP² in several studies (HD Natural History Study [NCT03664804], open-label extension of the tominersen Phase I/IIa study [NCT03342053], GENERATION HD1 [NCT03761849] and the Digital-HD study), into a dataset of over 1,000 individuals with HD.
- This dataset was divided into a development \bullet (N=141) and a validation dataset (N=319), which were used to build the HDDMS and to evaluate its generalisability and measurement properties.
- The top features were selected based on lacksquarehigh sensitivity to change (STC), high intraclass correlation coefficient (ICC), and

HD-ISS Stage 2 (N=113) and Stage 3 (N=610). cUHDRS, SWR, SDMT and TFC were multiplied by -1 so all features have the same directionality of progression (higher scores indicate the worse outcomes).

cUHDRS and the HDDMS. The calculation is based on mean change estimate from the GENERATION HD1 PLB and the pooled SD from the Q16W and PLB. No data were shown for cUHDRS at Week 36 to improve readability.

The HDDMS and its features significantly and consistently correlate with clinical anchors.

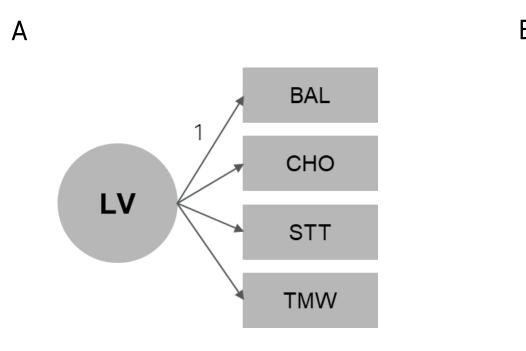
Figure 3. (A) Correlation heatmap; (B) relationship between HDDMS and cUHDRS



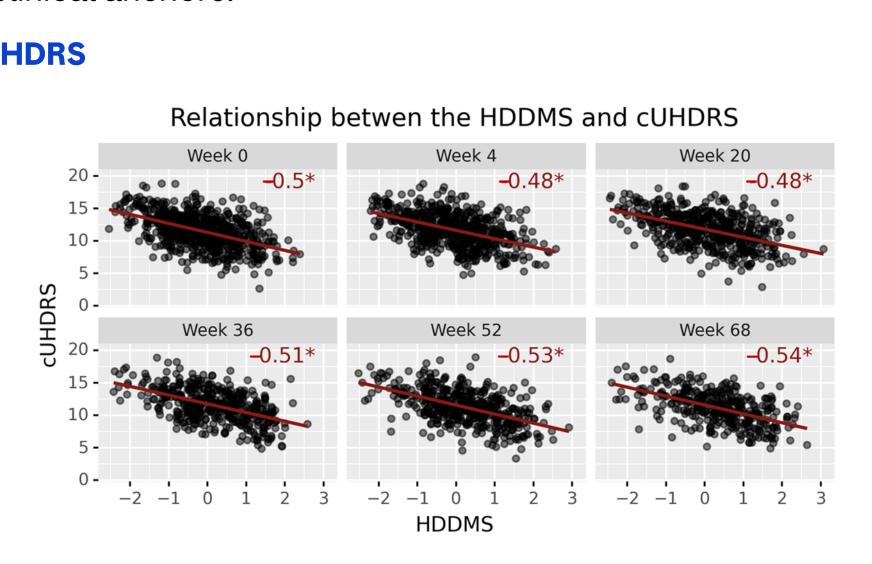
Label and colour = Pearson's r. * Indicates p < 0.001.

The HDDMS shows strong measurement invariance on the validation dataset.

Figure 4. (A) Measurement model; (B) fit indices of measurement model; (C) loadings and individual score weights



	Development	Validation	Full dataset	С	Feature	Loadings	Standardise loadings
alue led	0.72	0	0		BAL	1	0.6993
scaled	1	0.961	0.982		СНО	0.9159	0.7492
SEA led	0	0.062	0.043		STT	0.7830	0.6549
MR	0.063	0.060	0.031		тмw	0.7280	0.5580

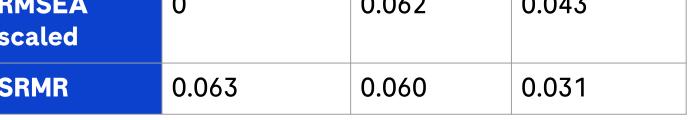


Relationship between the HDDMS and cUHDRS across weeks. The red line corresponds to a linear regression fit and the red number to the Pearson's r. * Indicates p<0.001. Data from GENERATION HD1 (N=730).

good cross-sectional correlation with cUHDRS. In addition, longitudinal feature progression had to be congruent with knowngroup differences from the Digital-HD study. STC was calculated from the estimates of change from baseline by a mixed model with repeated measures (MMRM) as: STC=estimate of change/standard deviation (SD).

The top-ranked feature of each test was input to factor analyses, combining both exploratory and confirmatory factor analysis, to find the best measurement model and to test measurement invariance over time and between datasets. Individual scores were calculated using the regression method.

Measurement model (with a loading of 1 for the BAL feature in order to identify the model). This model had the best fit on the development dataset compared with the other hypothesised measuremen models and was strongly invariant in regard to time (tested up to Week 68).



Fit indices of the measurement model with strong invariance over time (fixed intercept and fixed loadings). For validation, the same intercepts and loadings were used. Then, on the full dataset (development and GENERATION HD1), the model was re-fit (with strong measurement invariance over time) in order to fine tune model parameters later used to calculate individual scores.

RMSEA better be <0.08; SRMR, better be <0.08; CFI, better be >0.9. We used scaled test statistics to account for possible non-normality in the features.

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Abbreviations

BAL, balance; CFI, Comparative Fit Index; CHO, chorea; cUHDRS, composite Unified Huntington's Disease Rating Scale; DMP, digital monitoring platform; EQ-5D-5L, EuroQol 5-dimension 5-level; HD, Huntington's disease; HDDMS, HD digital motor score; HD-ISS, HD Integrated Staging System; ICC, Intraclass Correlation Coefficient; LV, latent variable; MMRM, mixed model for repeated measures; PLB, placebo; Q16W, every 16 weeks; RMSEA, root mean square error of approximation; SD, standard deviation; SDMT, Symbol Digit Modalities Test; SRMR, standardised root mean square residuals; STC, sensitivity to change; STT, Speeded Tapping Test; SWR, Stroop Word Reading; TFC, Total Functional Capacity; TMS, Total Motor Score; TMW, Two-minute Walking Test.

References

1. Schobel SA, et al. Neurology. 2017; 89:2495-2502; 2. Lipsmeier F, et al. J Med Internet Res. 2022; 24:e32997.

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