

Study protocol for the iMarkHD study in individuals with Huntington's disease

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Abstract

Background: Huntington's disease (HD) is still often defined by the onset of motor symptoms, inversely associated with the size of the CAG repeat expansion in the *huntingtin* gene. Although the cause of HD is known, much remains unknown about mechanisms underlying clinical symptom development, disease progression, and specific clinical subtypes/endophenotypes.

Objectives: In the iMarkHD study, we aim to investigate four discrete molecular positron emission tomography (PET) tracers and magnetic resonance imaging (MRI) markers as biomarkers for disease and symptom progression.

Methods: Following MRI optimisation in five healthy volunteers (cohort 1), we aim to recruit 108 participants of whom 72 are people with HD (PwHD) and 36 healthy volunteers (cohort 2). Pending interim analysis, these numbers could increase to 96 PwHD and 48 healthy controls. Participants will complete a total of 10 study visits, consisting of a screening visit followed by a clinical and MRI visit and PET visits at baseline, year 1, and year 2. PET targets include the cannabinoid 1, histamine 3, and serotonin 2A receptors, and phosphodiesterase 10A, whereas MRI will be multimodal, including, but not limited to, the assessment of cerebral blood flow, functional connectivity, and brain iron.

Results: Recruitment is currently active and started in September 2022.

Conclusions: By combining PET and multi-modal MRI assessments we expect to provide a comprehensive examination of the molecular, functional, and structural framework of HD progression. As such, the iMarkHD study will provide a solid base for the identification of treatment targets and novel outcome measures for future clinical trials.

Keywords: Huntington's disease, MRI, PET, neuroimaging, progression, biomarker.

Introduction

Huntington's disease (HD) is still often defined by the onset of motor symptoms, inversely associated with the size of the CAG repeat expansion in the *huntingtin* (*HTT*) gene.¹ Nevertheless, subclinical changes and pathological processes precede the initiation of clinical symptoms, often by many years.²⁻⁴ The availability of genetic testing and full penetrance of the *HTT* mutation in people with ≥ 40 CAG expansions provide a unique window of opportunity to examine the pattern of signs, symptoms, and neurobiological changes as they emerge, and study the clinical course of HD before the development of overt symptoms.⁵ Although the cause of HD is known, much remains unknown about the mechanisms underlying the development of clinical symptoms, disease progression, and specific clinical subtypes/endophenotypes. Moreover, the identification of easily obtainable, reliable, and robust biomarkers of HD progression is crucial for the development and evaluation of disease-modifying treatments.

Previous efforts to identify such biomarkers include the TRACK-HD study, which was a longitudinal study focusing on sensitive and reliable biomarkers with extensive annual assessments involving magnetic resonance imaging (MRI) and clinical measures in both premanifest and manifest people with HD (PwHD).^{2-4,6} In addition to motor progression, clear progression in cognitive dysfunction was observed,² particularly in relation to progression in apathy. Moreover, baseline apathy scores were significant predictors of later functional decline, as well as other neuropsychiatric symptoms associated with frontal lobe function, including affect and irritability.² MRI measures showed significant total brain volume reduction, most prominent in caudate and putamen volume, as well as striatal, corpus callosum and posterior white matter tract changes, with a strong relation to total functional capacity decline.³

Although MRI provides useful information about structural and functional organisation of brain changes in HD, including brain volume, perfusion, iron levels, and structural and functional connectivity, this technique is less useful to assess changes in specific neurotransmitters. For the latter, positron emission tomography (PET) may be deployed, aiding in understanding the progression and complexity of HD.⁷ Especially in relation to PET markers, iMarkHD will provide innovative longitudinal assessments of HD pathology, complemented by MRI imaging.

Methods

Protocol Title

Longitudinal Adaptive Study of Molecular Pathology and Neuronal Networks in People with Huntington's disease (PwHD) and Healthy Controls using Positron Emission Tomography (PET) and Multi-modal Magnetic Resonance Imaging (MRI). The short title for the project is: iMarkHD In Vivo Longitudinal Imaging of HD Pathology.

Study aims and objectives

The primary objectives of the iMarkHD study are to investigate the change in binding profile of four discrete molecular PET tracers across different disease stages in HD and age- and sex-matched healthy volunteers, measured over two years, to determine whether these PET imaging methods could be used as markers of HD progression and future treatment response in therapeutic trials. The PET tracers used in the study are: 1) [¹¹C]MDL100907 for serotonin receptor 2A (5HT2A);⁸ 2) [¹¹C]MK8278 for histamine 3 receptor (H3R);⁹ 3) [¹¹C]MePPeP for cannabinoid 1 receptor (CB1R);¹⁰ and 4) [¹¹C]IMA107 for phosphodiesterase 10A (PDE10A).¹¹

Secondary objectives include:

- 1) Quantitative cross-sectional and longitudinal assessment of brain structure, function, blood flow and iron level changes measured through MRI including arterial spin labelling (ASL), quantitative susceptibility mapping (QSM), quantitative relaxation times, diffusion tensor imaging (DTI) and resting-state functional MRI (rs-fMRI).
- 2) To explore the cross-sectional and longitudinal relationships between PET imaging measures and MRI measures with:
 - a. Number of *HTT* amino acid glutamine (CAG) repeats.
 - b. CAG-Age Product (CAP) score (=age x (CAG-30)/6.49).¹²
 - c. Clinical rating scales, including Unified Huntington's Disease Rating Scale (UHDRS), UHDRS Total Motor Score (UHDRS-TMS), UHDRS Total Functional Capacity (UHDRS-TFC) scores, and psychiatric and cognitive assessments.
 - d. Disease duration as assessed by time at which first motor clinical manifestation was noticeable.
- 3) To develop and evaluate the application of combined multi-modal MRI and PET in assessing cross-sectional and longitudinal structural, functional, and molecular changes in the brain of PwHD compared to healthy controls.
- 4) To develop probabilistic tractography methodology fusing combined PET/DTI for assessing pathway-based functional molecular changes and associations in line with known brain circuits and primary and secondary objectives in PwHD and healthy controls.
- 5) To determine whether the [11C]IMA107 PET marker can be a prognostic marker of HD phenoconversion (clinical motor diagnosis) and how it compares to the prognostic accuracy of the other three PET markers included in the study.

Rationale for positron emission tomography targets

PDE10A activity appears to be one of the most promising markers for disease progression as PET imaging data in PwHD have shown that changes in PDE10A activity in the striatum may be one of the first detectable neuroimaging changes in premanifest PwHD.¹³ Similarly, striatal PDE10A mRNA and protein levels are reduced in HD transgenic mice prior to the onset of motor symptoms.^{14,15} Due to these early changes occurring in premanifest PwHD,^{16,17} it has been suggested that PDE10A activity could serve as a marker to predict the occurrence of manifest motor symptoms and could be more sensitive than the dopamine-receptor and volumetric methods currently used.¹⁷

CB1R has been implicated as neuroprotective through an effect on glutamatergic synapses. In animal models of striatal damage, activation of CB1R on cortico-striatal projections selectively protect medium spiny neuron populations.¹⁸ Interestingly, genetic deficiency of CB1R has been shown to worsen disease signs in the N171-82Q transgenic HD mice,¹⁹ and a delayed onset of HD-like symptoms in other rodent HD models were associated with less pronounced loss of this receptor.²⁰ Similar findings were obtained in the R6/2 HD mice.¹⁴ In PwHD, small-scale PET studies have shown profound early and widespread reduction of CB1R availability.²¹

Serotonergic neurotransmission is an area of interest across several neurodegenerative conditions, including HD. Decreased serotonergic neurotransmission has been linked to depressive symptoms, at least in the R6/1 HD mouse model.²² Moreover, specifically 5HT2A shows an age-dependent decrease in the striatum, caudate, and hippocampus of zQ175 HD mice,⁸ in line with findings from the R6/1 HD mice.²³ In the zQ175 HD mouse model, a PET study has shown that at six months of age, striatal binding ratios for [¹¹C]-MDL 100907, as a marker for 5HT2A, was decreased by 11% compared to wild-type mice. In addition, an overall reduction in hippocampal and cortical 5HT2A was observed with further reductions of striatal binding at nine months of age, whereas PDE10A

reached a plateau.⁸ This suggest that 5HT2A might be a useful biomarker in more advanced stages of HD.

Finally, the central histaminergic neuropeptidergic system also shows promise as a biomarker for HD progression. Histamine has been suggested to play a role in the neuronal and oxidative damages observed in HD mouse models, focusing on striatal changes.²⁴ Evidence from PwHD shows not only an increase and a diurnal phase shift in L-histidine expression as a measure for histamine production in the hypothalamic tuberomammillary nucleus, but also a selective increase in H1, H2, and H3 histamine receptor expression in cortical areas,^{25,26} indicating hyperactivity of the neuronal histaminergic system. This is likely to be of relevance given the effect of histamine on neuronal and oxidative damage which has been observed in quinolinic acid induced HD mouse models.²⁵ Another animal study has demonstrated that overactivation of dopamine D1 receptors leads to an imbalance in dopaminergic neurotransmission and cell death, which could be mitigated by targeting dopamine D1 and H3 receptor complexes through an H3R antagonist. Moreover, receptor complex levels were associated with disease duration suggesting a possible role as biomarker for H3R.²⁷

Rationale for multimodal magnetic resonance imaging

Conventional T1-weighted structural MRI, at a spatial resolution in all three dimensions of 1 mm, allows the quantification of total white and grey matter as well as that of relevant anatomical regions of interest such as the caudate and striatum relevant to HD pathology,²⁸ in addition to these data being used for co-registration with PET imaging data to allow anatomical delineation of the receptor distribution. ASL will provide quantitative maps of cerebral perfusion and blood flow, which have been previously shown to be locally reduced in PwHD not solely explained by tissue atrophy.²⁹ The rs-fMRI will be used to evaluate functional connectivity through measures of blood-oxygen- level

dependent (BOLD) signal as there is some evidence to suggest that functional connectivity may be profoundly altered in PwHD.³⁰ Emerging evidence suggests that dynamic functional connectivity metrics may indicate changes in macroscopic neural activity patterns underlying critical aspects of cognition and behaviour.³¹ DTI, used to probe the integrity of structural white matter connections, has previously been used to show that diffusion metrics were significantly altered in premanifest and early manifest PwHD.^{32,33} Finally, QSM, sensitive to cerebral iron accumulation, has also been previously shown to be abnormal in PwHD. For example, iron deposition in subcortical areas is increased in premanifest and manifest PwHD.³⁴ The iMarkHD study will evaluate which MRI measure or combination thereof is most predictive of disease progression.

Ethical approval

The study received ethical approval from London - Bromley Research Ethics Committee (reference 19/LO/0339; Integrated Research Application System reference 242859). This study will be conducted in accordance with the principles of the Declaration of Helsinki and General Data Protection Regulation. Written informed consent will be obtained from all participants prior to study-related activities.

Study design

The iMarkHD study is a UK-based, longitudinal, adaptive, observational, multi-target PET molecular and multi-modal MR imaging study designed to identify candidate disease progression and treatment response biomarkers to increase knowledge of HD-related mechanisms. The study is registered on clinicaltrials.gov (<https://classic.clinicaltrials.gov/ct2/show/NCT03434548>). The recruitment target for the study is 113 participants across two cohorts. Cohort 1 consists of five healthy participants

undergoing MR scanning to optimise data acquisition and MR sequences for Cohort 2. In Cohort 2, 72 PwHD and 36 healthy volunteers across four different groups (Table 1) will be recruited:

- Group B: PwHD without symptoms (approximately HD-ISS stage 0/1; ³⁵ CAP score >70) (n=24)
- Group C: PwHD with early disease (approximately HD-ISS stage 2) ³⁵ (n=24)
- Group D: PwHD with later disease (approximately HD-ISS stage 3) ³⁵ (n=24)
- Group E: Healthy volunteers (n=36)

Table 1 describes the eligibility criteria for the study and for each of the groups. We have chosen to allow participants with a TFC score of 12 to participate in Groups A (see below) and B and not consider a score of 12, in the absence of other major symptoms, indicative of more advanced stages of HD. An example where a participant in these groups scores 12 on the TFC is unemployment which can occur without clinical signs or symptoms. This is in line with previous neuroimaging studies in PwHD and has also been taken into account in the HD-ISS classification. ³⁸

Pending interim analysis taking place at 25-50% completion of baseline assessments of groups B and E, a decision will be taken whether or not to introduce a Group A consisting of PwHD without symptoms and expected far to onset of motor symptoms (CAP score ≤ 70 ; approximately HD-ISS stage 0) ³⁵ (n=24), as well as 12 additional healthy volunteers. The decision will be based on whether a significant marker difference between PwHD in Group B and healthy controls is present, supporting the rationale for investigating the earliest change in marker profile in further to expected motor onset PwHD (Group A). Phenoconversion in the iMarkHD study, separating participants in Group C from those in Group B, is defined by the occurrence of motor symptoms (TMS scores ≥ 7) and/or decline in total functional capacity (TFC scores 11 and 12). In those Group C participants in whom higher motor scores may be present (TMS 24-33) a further requirement is a score of >50 on

the symbol digit modalities test (SDMT) which has been suggested as a useful early neuropsychological sign to indicate disease process status prior to reaching motor diagnosis criteria.

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In cohort 1, the five healthy volunteers will undergo a screening visit to determine eligibility, followed by two visits for MR and MR retest scanning with no further visits. For Cohort 2, following a screening visit to determine eligibility, participants will undergo clinical, PET and MR imaging assessments at baseline, and 1- and 2-year follow-up (Figure 1). Study procedures for cohort 2 are outlined below.

Study sample and recruitment

Participants for this study will be recruited through King's College London and King's College Hospital, London, United Kingdom. In addition to local recruitment, PwHD will be referred by centres specialised in HD, currently consisting of 11 participant identification centres (PICs) from academic healthcare and National Health Service (NHS) organisations across the United Kingdom. The referring PICs and the clinical staff at King's College London will identify potentially eligible participants and their willingness to participate in the study. Recruitment will be supported through a public advertisement approved by the Health Research Authority (HRA). All participants will receive travel reimbursement as well as HRA-approved stipends for each attended study visit. Further reimbursement will be provided for overnight stay, where applicable, for participants travelling from outside London.

Sample size

The sample size of each group in the iMarkHD study has been based on previously reported differences in cross-sectional studies showing standardised differences (Cohen's d) between PwHD and healthy volunteers in PET markers to range between 1.8 ([¹¹C]MePPEP PET in the striatum) and 2.55 ([¹¹C]MePPEP PET in cortex) (unpublished data), in addition to differences between 0.5 ([¹¹C]IMA107 in substantia nigra), and 2.56 ([¹¹C]IMA107 in the putamen).³⁷ Based on this, 20 participants per group (100 in total), will provide 80% power to detect baseline differences between PwHD subgroups and healthy volunteers corresponding to a Cohen's d of 0.78 or larger with a one-sided test, 5% type I error rate. In order to ensure that 20 participants reach 2-year follow up, 24 participants will be recruited per group, to account for the expected approximate 20% dropout rate (10% attrition per year). Differences across groups will only be tested when differences versus healthy volunteers are found.

Power to detect longitudinal differences has been estimated for a mixed-model with repeated measures; one baseline and two follow up measurements, with a general correlation structure (assuming 0.5 between baseline and 2-years, and 0.8 between consecutive measurements). Based on 20 participants with complete follow up this will provide 80% power to detect standardized differences of 0.91 or higher, with a 2-sided 5% type I error rate, between PwHD participants and healthy volunteers. A previous study based on a small number of participants (n=12) has indeed shown that correlations between PDE-10A expression and probability for symptomatic conversion after 2 years, in far from motor onset PwHD, range from 0.59 in the higher motor-thalamic-nuclei/striatal, to 0.85 in the higher motor-thalamic-nuclei/striatopallidal.³⁷ It is expected that PET markers under investigation in this study will be more sensitive and detect larger changes earlier than previously established clinical and MRI markers.^{2-4,6,32,33}

308 A sample size of 36 healthy volunteers has been chosen to adequately age match against three PwHD
309 groups based on previous data in similar groups in the TRACK-HD study at visit 1.²⁻⁴

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311 *Study procedures*

312 For all participants in the study a screening visit will be performed to assess eligibility and to enable
313 grouping of participants based on the measures outlined in Table 1. During the screening visit
314 participants will be consented to the study and demographics, such as age and sex, will be collected.
315 In addition, past medical history will be obtained along with current medication use, and a medical
316 and neurological examination performed, including Snellen chart and Rinne and Weber testing. Also,
317 vital signs will be recorded and an electrocardiogram performed. To check for exclusion criteria,
318 blood and urine samples will be obtained along with analysis of *HTT* CAG-repeat length that will be
319 anonymously linked to the Enroll-HD database,³⁸ as well as performing the Structured Clinical
320 Interview for DSM-5 Research Version (SCID-5-RV). Finally, to group participants into one of the
321 iMarkHD groups, the Unified Huntington's Disease Rating Scale (UHDRS)³⁹ Total Motor Score
322 (UHDRS-TMS), Diagnostic Confidence Level (UHDRS-DCL), and Total Functional Capacity
323 (UHDRS-TFC) will be assessed, along with Symbol Digit Modalities Test (SDMT),⁴⁰ Hospital
324 Anxiety and Depression Scale (HADS),⁴¹ and Mini Mental State Examination (MMSE) scores.⁴²

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326 As mentioned above, for Cohort 1 the five healthy volunteers will undergo a screening visit, followed
327 by a MR visit and a MR retest visit. These two latter visits will consist of MR scanning taking up to
328 two hours per visit. MRI scanning will be performed on a GE MR750 3Tesla MRI scanner (for
329 Cohort 1 and 2). In Cohort 2, all participants will complete a total of 10 study visits, consisting of a
330 screening visit (visit 1), followed by a combined clinical and MR visit (visit 2) and two PET visits
331 (visits 3 and 4; during each PET visit participants will undergo two PET scans), at baseline, year 1

(visits 5, 6 and 7), and year 2 (visits 8, 9, and 10). An overview of study visits is provided in Figure 1 and clinical outcomes are summarised in Table 2.

PET targets and MR sequences that will be performed are summarised in Figures 2 and 3 and clinical assessments are summarised in Table 2. For Visits 2, 5 and 8, an effort will be made to perform clinical assessments and the MR sessions in the same visit, but based on availability it is possible that these visits may be performed over two separate days; for Visits 3, 4, 6, 7, 9 and 10, an effort will be made to perform two PET scans on the same day, but based on availability it is possible that these visits may be performed over two separate days. The overall study design is summarised in figure 1.

For Cohort 1 the first MR visit will be performed within 30 days of screening, and the second visit will be performed within 14 days of the first MRI visit. For Cohort 2, visit V2 will be performed within 40 days of screening (V1), and the PET visit V3 will be performed within 30 days from V2 with V4 taking place within 14 days from V3. The same timelines (i.e. first PET visit within 30 days of MRI, and clinical visit and second PET visit within 14 days of first PET visit) will be repeated for year 1 and year 2 follow-up.

Data analysis

All PET and MRI outcome markers will be analysed using a generalised linear model, with the absolute values of each marker as dependent variable, and subgroup categorical variables as independent, in addition to sex and age as covariates in the model. Benjamini–Hochberg false discovery rate correction at 0.1 to correct for multiple testing will be used. Marginal means, and standardised effect sizes will be used to report differences across groups. In addition, partial correlations across different markers and clinical measurements, correcting for age and sex, will be

used. To analyse longitudinal differences across groups fit linear mixed models for repeated measures, with random intercept and slope, will be used. The presence of non-linear changes will be tested by adding a quadratic term for time. The outcome variable will consist of the absolute levels of the individual PET and MRI markers at baseline and the two annual follow up measurements, and categorical subgroups as the main independent variable.

Data from clinical assessments, including rating scales to measure disease progression, demographics, and other baseline characteristics, will be summarised using descriptive statistics and linked to PET and MRI markers using mixed linear models and correlations, where appropriate.

Data handling and storage

All data will be handled confidentially and processed, stored, and disposed of in accordance with the General Data Protection Regulation and all applicable legal and regulatory requirements in the UK. Data collection will be performed with the use of paper and secured electronic case report forms (CRF). The database will only be accessible for authorised personnel via a unique login and user ID. All participant data will be coded, and any identifiable data removed prior to uploading. All paper source documents will be stored in a secure location for the duration of the study and any time following this depending on legal and regulatory requirements.

PET neuroimaging data will initially be stored on local servers at the PET imaging acquisition centre and within one week after image acquisition securely transferred via Microsoft One Drive to the department of neuroimaging at King's College London. Here, together with the MRI neuroimaging data, the data will be securely stored on internal servers. At the end of the study, these data will also be transferred via secure environments to CHDI.

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381 *Benefits and risks assessment*

382 As this is a non-interventional study, no excessive risks or ethical concerns are anticipated. Any
383 adverse events will likely be restricted to blood sampling and MRI and PET procedures and will be
384 recorded in the paper and electronic CRFs. In relation to PET imaging the main risk to participants
385 consists of the risk due to ionising radiation exposure from injected radioligands and associated low
386 dose computed tomography (CT). The target dose for the individual PET tracers is 200 MBq
387 (between 0.76 and 1.08 mSv per scan) in addition to the effective radiation dose associated with CT
388 scanning (0.36 mSv per scan). As such, it has been calculated that the total effective dose for each
389 participant is 5.08 mSv per year and 15.20 mSv in total across all study visits. This compares to the
390 average 2.3 mSv of annual background radiation for UK residents. The radiation dose has been
391 chosen as providing acceptable image quality while minimising participant risk. While any additional
392 ionising radiation exposure may increase the risk of future malignancy, this risk is mitigated by
393 clinical safety monitoring and safety assessment procedures during study participation and
394 inclusion/exclusion criteria and restrictions applied during study participation. As there are no
395 alternative methods to PET targets in our study, we feel the added risk due to the additional radiation
396 exposure is justified.

397

398 Serious adverse events will also be reported to the study sponsor. Planning of study visits will be
399 done depending on participants' availability and wishes, insofar as possible. The frequency and
400 duration of the scanning sessions might cause some distress to participants, but this will be
401 specifically assessed as part of the eligibility criteria. All researchers involved in this study are
402 experienced in neurological clinical research. Further potential untoward effects will be minimised by
403 pilot testing of MRI sequences in Cohort 1, consisting of healthy volunteers.

404

405 **Discussion**

406 The aim of the iMarkHD study is to evaluate how specific molecular, functional, and structural
407 changes may influence the development of symptoms in PwHD and how this might aid in the
408 development of imaging biomarkers that can characterise and predict events over the course of the
409 disease. Ultimately, the outcomes of this study may be used as outcome measures in future clinical
410 trials and identification of novel treatment targets. PET molecular imaging is a powerful in vivo tool
411 to investigate brain function such as metabolism, and receptor and enzyme distributions. As an
412 analytical imaging method, PET has the potential to give both structural and kinetic information and,
413 in comparison with other imaging techniques with high sensitivity and temporal resolution.⁴³ The
414 PET assessment will be complemented by MRI to provide information on structural and functional
415 organisation of the brain in PwHD, as well as brain volume, perfusion, iron levels, and structural and
416 functional connectivity. The combination of these PET and MRI techniques have the potential to
417 significantly contribute to our understanding of the complexity of HD.⁷

418

419 As with any study, some challenges are expected to arise during the implementation process of the
420 iMarkHD study. These include participant recruitment in a single centre study, participant retention
421 in relation to the relatively high number of imaging sessions, as well as factors related to eligibility
422 criteria and the inclusion of PwHD without symptoms (HD-ISS stages 0-1). We aim to address these
423 challenges by actively engaging with PICs to maximise our recruitment area and to engage to the
424 largest extent possible with PwHD across the UK; currently, we are working with 11 PICs in the UK
425 with a large geographical distribution and catchment area. This will enable us to screen a large
426 population of PwHD, offsetting some of the stricter eligibility criteria in the study protocol.
427 Moreover, as the HD community is generally very motivated with high willingness to participate in

research, we expect to overcome the challenges in relation to recruitment with accrual of sufficient participants for each group. Another possible challenge is the adequate classification of each participant into the relevant groups described in the iMarkHD study protocol. Rather than relying on what is considered as increasingly outdated, premanifest and manifest HD,³⁵ we have included a more dynamic classification system based on TFC and TMS, in addition to CAP scores, similar to designs used in other HD studies.⁴⁴ This system will allow for more accurate classification of participants and reduce subjective rater classification into premanifest and manifest.³⁵

The unique aspect of the iMarkHD study is mainly related to its multimodal assessment of possible HD markers. Since it is likely that multiple pathophysiological changes influence disease progression over time, combining PET and multi-modal MRI assessments in this study will likely provide a comprehensive examination of the molecular, functional, and structural framework of HD progression in the brain. In addition, the inclusion of a group of healthy control volunteers to compare PET and MRI measurements at different disease stages will further contribute to our understanding of the development of symptoms and disease progression in HD. As such, the study will provide a solid base for the identification of treatment targets and could provide novel outcome measures for future clinical trials evaluating new targeted therapies. PET imaging offers molecular insights crucial for understanding disease mechanisms and progression, while complementing other imaging modalities such as MRI. In addition, developments over the years have resulted in safer radiotracer development utilising lower effective doses. Therefore, the use of PET in future clinical trials and other clinical research outweigh the risks when used responsibly.

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Conflict of interest

The authors declare that the research described is conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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590 **Table 1. Eligibility criteria per group for the iMarkHD study.** Abbreviations: PwHD: people with
591 Huntington's disease; TMS: Unified Huntington's Disease Rating Scale (UHDRS) Total Motor
592 Score; TFC: UHDRS Total Functional Capacity; CAP: CAG Age Product; SDMT: Symbol Digit
593 Modalities Test; HD: Huntington's disease.

Inclusion criteria for the entire cohort 1. Age ≥ 21 to ≤ 75 years at the time of screening 2. Adequate visual (Snellen chart) and auditory (Rinne and Weber tests) acuity to complete the psychological testing as determined by the investigator 3. Capability of giving informed consent 4. Willing to comply with highly effective contraceptive measures 5. Vital signs (after three minutes resting in a supine position) which are within the following ranges at the screening evaluation: a. Systolic blood pressure 85-160 mmHg b. Diastolic blood pressure 50-100 mmHg c. Heart rate 45-110 bpm 6. Absence of clinically significant diseases (excluding disease under study) or clinically significant abnormal laboratory values 7. Ability to travel to imaging and clinical assessment centres in London
Group A (PwHD without symptoms far from expected motor onset) 1. PwHD with ≥ 40 CAG repeats in the <i>Htt</i> gene 2. TMS ≤ 6 AND TFC ≥ 12 AND CAP ≤ 70
Group B (PwHD without symptoms) 1. PwHD with ≥ 40 CAG repeats in the <i>Htt</i> gene 2. TMS ≤ 6 AND TFC ≥ 12 AND CAP > 70
Group C (PwHD with early disease) 1. PwHD with ≥ 40 CAG repeats in the <i>Htt</i> gene 2. If one of the following criteria is met a. TMS ≤ 6 AND TFC = 11 b. TMS is between 7 and 23 inclusive AND TFC is between 11 and 13 inclusive c. TMS is between 24 and 33 inclusive AND SDMT > 50 AND TFC is between 11 and 13 inclusive
Group D (PwHD with later disease) 1. PwHD with ≥ 40 CAG repeats in the <i>Htt</i> gene 2. If one of the following criteria is met: a. TMS ≤ 6 AND TFC is between 7 and 10 inclusive b. TMS is between 7 and 23 inclusive AND TFC is between 8 and 10 inclusive c. TMS is between 24 and 33 inclusive AND SDMT > 50 AND TFC is between 7 and 10 inclusive d. TMS is between 7 and 23 inclusive AND TFC = 7 e. TMS > 23 AND SDMT ≤ 50 AND TFC is between 7 and 13 inclusive f. TMS > 33 and SDMT > 50 AND TFC is between 7 and 13 inclusive
Group E (Healthy controls) 1. Age (± 8 years) and sex matched with PwHD

2. No known family history of HD or have known family history of HD but have been tested for the huntingtin gene glutamine codon expansion and are not at genetic risk for HD (CAG < 36)

Exclusion criteria for the entire cohort

1. Presence or history of other neurological condition likely to interfere with imaging or abnormal neurologic examination findings suggestive of a central nervous system pathology (other than HD)
2. Presence of history of primary psychiatric disorders unrelated to HD
3. Participants using any medications with known actions on any of the PET targets
4. Pregnancy confirmed by a positive urine pregnancy test
5. Participants who are currently breastfeeding or intend to breastfeed during the study
6. Contraindication to MRI
7. History of alcoholism or substance abuse within three years prior to study entry
8. Failure of drug screen for substances of abuse such as amphetamines, barbiturates, benzodiazepines, methadone, opiates, cocaine, cannabinoids, phencyclidine, and creatine
9. History of cancer
10. Claustrophobia
11. Significant back pain that makes prolonged laying on the PET or MRI scanner intolerable
12. Contraindication for arterial cannulation as judged by the Allen test and the laboratory blood screening for coagulopathy
13. Inability to communicate or cooperate with the Principal Investigator/iMarkHD Team for any reason
14. Participants who are enrolled in or participated in clinical trials testing the efficacy of novel therapeutics with action on the specific PET targets being tested within 3 months of screening
15. Any concurrent conditions that could interfere with the safety and/or tolerability measurements.

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606 **Table 2. Clinical assessments in cohort 2 of the iMarkHD study.** Abbreviations: UHDRS: Unified
607 Huntington’s Disease Rating Scale; TMS: Total Motor Score; DCL: Diagnostic Confidence Level;
608 TFC: Total Functional Capacity; FAS: Functional Assessment Scale; IS: Independence Scale;
609 HDQoL: Huntington’s Disease Quality of Life; HD: Huntington’s disease.

Motor assessments UHDRS-TMS UHDRS-DCL	Cognitive assessments Cognitive Assessment Battery: • Symbol Digit Modalities Test • Emotion Recognition • Trail Making Test • Hopkins Verbal Learning Test Revised • Paced Tapping Test • Tower-Z task Stroop Colour and Word Reading Test and Interference Test Categorical Verbal Fluency Test Mini-Mental State Examination
Functional assessments UHDRS-TFC UHDRS-FAS UHDRS-IS Physical Performance Test HDQoL	
Neuropsychiatric assessment Problem Behaviour Assessment Short Version Hospital Anxiety and Depression Scale Apathy Evaluation Scale HD Clinical Characteristics Assessment	

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Figure 1. Overall study design of the iMarkHD study. Abbreviations: MRI: magnetic resonance imaging; PET: positron emission tomography; PwHD: people with Huntington’s disease.

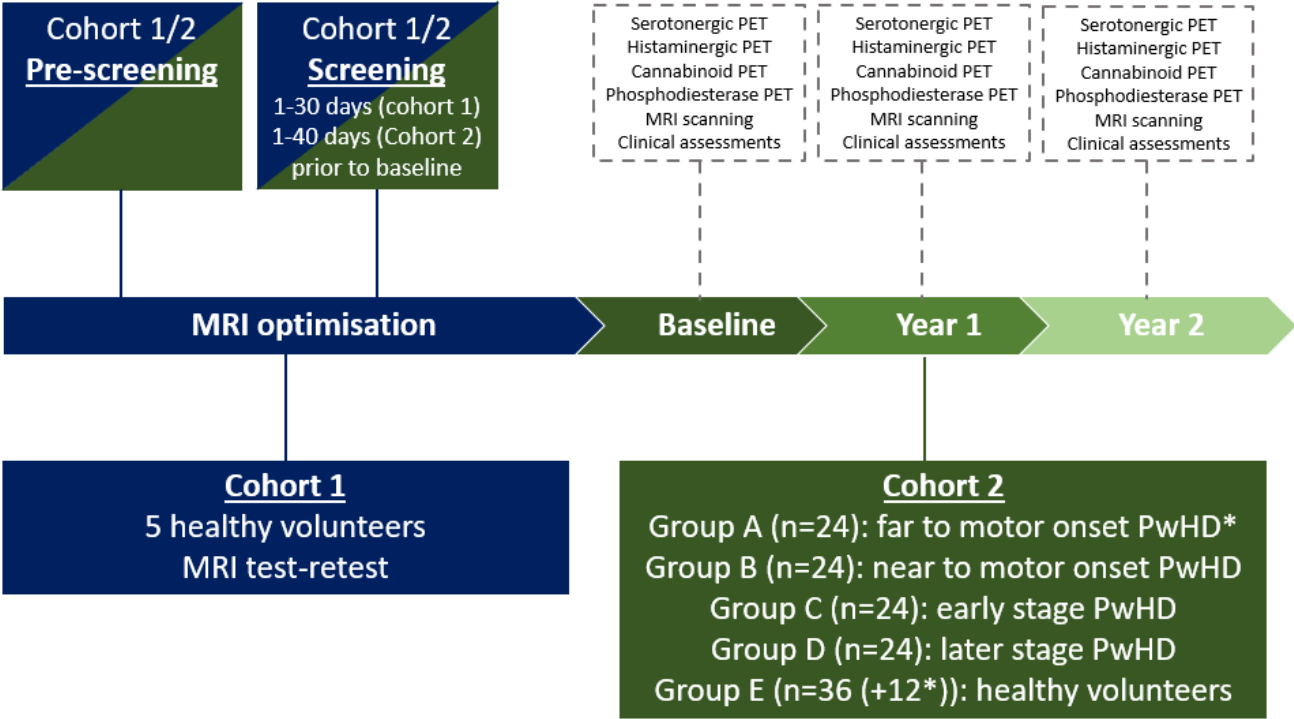


Figure 2. Positron Emission Tomography outcomes for the iMarkHD study. The figure shows receptors (centre panel) and PET tracers (bottom panel) with illustrative scan results. Abbreviations: 5HT_{2A}R: Serotonin receptor 2A; H₃R: Histamine 3 receptor; CB₁R: Cannabinoid 1 receptor; PDE10A: phosphodiesterase 10A enzyme.

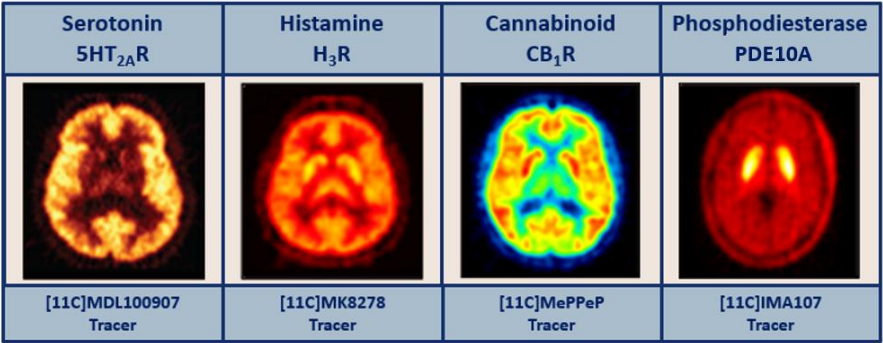
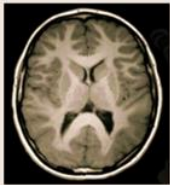
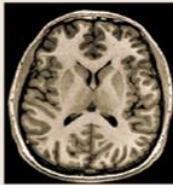
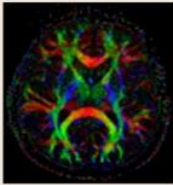
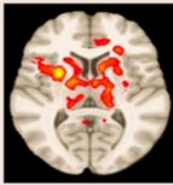
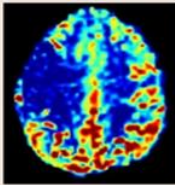
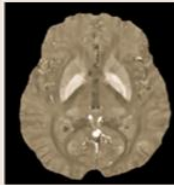


Figure 3. Magnetic resonance imaging outcomes for the iMarkHD study. The figure shows the different magnetic resonance imaging (MRI) outcomes (top panel) and what these outcomes represent (bottom panel) alongside illustrative scan results. Abbreviations: MP-RAGE: Magnetisation prepared rapid acquisition gradient echo; MPM: Multi-Parametric Mapping; DTI: Diffusion Tensor Imaging; fMRI: functional MRI; ASL: Arterial Spin Labelling; QSM: Quantitative Susceptibility Mapping.

MP-RAGE	MPM	DTI	fMRI	ASL	QSM
					
Structural & PET registration and analysis	T1 and T2 map quantification	White matter structures	Functional connectivity	Cerebral blood flow	Brain iron levels