Supplemental Methods

Participants

Supplementary Table 1 displays the demographic data for the full sample reported in Scahill et al. (2020) [1].

Supplementary Table 1. Demographics.

	Premanifest HD (N = 64) Healthy Controls (N = 67) P value			
Age	29.0 (5.60)	29.10 (5.70)	.95	
Sex	53% Females (34)	57% Females (38)	.81	
IQ	102.40 (7.50)	103.50 (8.30)	.42	
Years to onset	23.60 (5.80)	N/A		

Participants were identified and recruited from the Enroll-HD study https://www.enroll-hd.org/, regional Genetic and HD centres across the UK who were established as patient identification sites, and through broader efforts such as via the Huntington's Disease Association https://www.hda.org.uk/ and the Huntington's Disease Youth Organisation https://hdyo.org/.

Eligibility screening

Prior to enrolment, each participant was interviewed to determine whether they meet the eligibility criteria below. For gene carriers, CAG expansion in the HD gene was confirmed by obtaining the genetic report from an accredited laboratory. This was to confirm CAG expansion and standardise CAG sizing for statistical analysis.

Inclusion criteria

- a. Are 18-40 years of age, inclusive; and
- b. Are capable of providing informed consent and
- c. Are capable of complying with study procedures and

For the **Healthy Control** group, participants eligible are persons who meet the following criteria:

- d. Have no known family history of HD (family control or community control); or
- e. Have known family history of HD but have been tested for the huntingtin gene CAG expansion and are not at genetic risk for HD (CAG < 36*) (gene negative).

For the **Young Adult Premanifest HD** group, participants eligible are persons who meet the additional following criteria:

- f. Do not have clinical diagnostic motor features of HD, defined as Unified Huntington's
 Disease Rating Scale (UHDRS) Diagnostic Confidence Score [2] < 4; and
- g. Have CAG expansion ≥ 40*; and
- h. A disease burden score (DBS) [3] ≤ 240**2

The rationale for this DBS cut-off is that this boundary corresponds approximately to >18 years to estimated disease onset according to the Langbehn formula [4].

Exclusion criteria

- a. Current use of investigational drugs or participation in a clinical drug trial within 30 days prior to study visit; or
- b. Current intoxication, drug or alcohol abuse or dependence; or
- c. If using any antidepressant, psychoactive, psychotropic or other medications or nutraceuticals used to treat HD, the use of inappropriate (e.g., non-therapeutically high) or unstable dose within 30 days prior to study visit; or
- d. Significant medical, neurological or psychiatric co-morbidity likely, in the judgment of the Principal Investigator, to impair participant's ability to complete essential study procedures;
 or
- e. Predictable non-compliance as assessed by the Principal Investigator; or
- f. Inability or unwillingness to undertake any of the essential study procedures; or
- g. Needle phobia; or

- h. Contraindication to MRI, including, but not limited to, MR-incompatible pacemakers, recent metallic implants, foreign body in the eye or other indications, as assessed by a standard pre-MRI questionnaire; or
- i. Pregnant (as confirmed by urine pregnancy test); or
- j. Claustrophobia, or any other condition that would make the subject incapable of undergoing an MRI.

For the optional CSF collection only

- k. Needle phobia, frequent headache, significant lower spinal deformity or major surgery; or
- I. Antiplatelet or anticoagulant therapy within the 14 days prior to sampling visit, including but not limited to: aspirin, clopidogrel, dipyridamole, warfarin, dabigatran, rivaroxaban and apixaban; or
- m. Clotting or bruising disorder; or
- n. Screening blood test results outside the clinical laboratory's normal range for the following: white cell count, neutrophil count, lymphocyte count, haemoglobin (Hb), platelets, prothrombin time (PT) or activated partial thromboplastin time (APTT); or
- o. Screening blood test results for C-reactive protein (CRP)>2× upper limit of normal; or
- p. Exclusion during history or physical examination, final decision to be made by thePrincipal Investigator; including but not limited to:

i any reason to suspect abnormal bleeding tendency, e.g. easy bruising, petechial rash; or ii any reason to suspect new focal neurological lesion, e.g. new headache, optic disc swelling, asymmetric focal long tract signs; or

iii any other reason that, in the clinical judgment of the operator or the Principal Investigator, it is felt that lumbar puncture is unsafe.

CANTAB Intra-Extra Dimensional Set Shift Task

The CANTAB IED[5] is a 7-minute test measuring cognitive flexibility and has similarities to a computerised version of the Wisconsin Card Sorting Test[6]. The task initially features rule acquisition and reversal learning and then attentional set formation and set shifting. Participants are presented with two artificial dimensions including pink shapes and white lines.

Through trial and error the participant must select the correct rule. After six correct responses, the stimuli and/or rule changes. There are nine stages to the test. Initially the test presents simple stimuli with just one dimension (pink shapes). These later change to compound stimuli (white lines overlaid on pink shapes). Early in the test the shifts are intra-dimensional (ID) (pink shapes are relevant) to establish set formation. This stage assesses generalisation of learning. Then at stage 8 a crucial extra-dimensional (ED) shift occurs, white lines become relevant (attentional set-shifting). This stage assesses cognitive flexibility. This latter stage is followed by a final reversal of the rule. Importantly, the task is not time-limited and participants have 50 trials to reach learning criterion. Outcomes measures for the present study include the number of pre-ED errors (stages 1-7) and the ED shift errors (stage 8). Pre-ED errors measure learning performance on the stages up to but not including the critical ED shifting stage.

Supplemental Results

Separate analysis of covariance (ANCOVAs), controlling for age, sex and IQ were conducted between pre-HD and HC groups to compare pre-ED latency and ED latency. The results showed that there was no statistically significant difference between pre-HD and HC groups on either the pre-ED latency (F(1,99)=.691, p=.408 \mathfrak{y}_p^2 =.007) or ED latency (F(1,99)=.463, p=.498 \mathfrak{y}_p^2 =.005).

Full results for the functional connectivity comparison between pre-HD and HC groups are displayed in Supplementary Table 2.

Supplementary Table 2. Functional connectivity group comparisons.

	Premanifest HD	Healthy Controls	t-value	p-value	\mathfrak{y}_p^2
Right Caudate - Right vIPFC	.165 (.342)	.072 (.339)	1.836	.179	.018
Right Caudate - Left vIPFC	.084 (.352)	.081 (.307)	.012	.912	.001
Right Caudate - Right dIPFC	.008 (.297)	.024 (.300)	.073	.788	.001
Right Caudate - Left dlPFC	.015 (.279)	.031 (.300)	.029	.864	.001
Left Caudate – Right vIPFC	.135 (.333)	.060 (.337)	1.358	.247	.014
Left Caudate- Left vIPFC	.075 (.325)	.054 (.302)	.140	.709	.001
Left Caudate - Right dIPFC	.028 (.273)	023 (.351)	.631	.429	.006
Left Caudate - Left dIPFC	.049 (.261)	.024 (.316)	.196	.659	.002
Right Ventral Striatum – Right vIPFC	.118 (.345)	.016 (.368)	1.693	.196	.017
Right Ventral Striatum - Left vIPFC	.096 (.363)	.074 (.357)	.040	.841	.001
Right Ventral Striatum - Right dIPFC	107 (.286)	093 (.353)	.077	.782	.001
Right Ventral Striatum - Left dIPFC	089 (.260)	105 (.261)	.022	.883	.001
Left Ventral Striatum - Right vIPFC	.088 (.334)	018 (.390)	1.663	.200	.017
Left Ventral Striatum - Left vIPFC	.127 (.359)	.064 (.400)	.460	.499	.005
Left Ventral Striatum - Right dIPFC	115 (.274)	119 (.355)	.006	.938	.001
Left Ventral Striatum - Left dIPFC	079 (.259)	093 (.279)	.030	.864	.001

The full results for the partial correlations, controlling for age, IQ and sex, between ED errors and pairwise functional connectivity for both the pre-HD and HC groups are displayed in Supplementary Table 3.

Supplementary Table 3. Correlations between all pairwise ROIs and ED Errors.

	Premanifest HD ED Errors		Healthy Control	
			ED Errors	
	R-Value	P-Value	R-Value	P-Value
Right Caudate – Right vIPFC	.066	.657	.351	.011
Right Caudate - Left vIPFC	243	.096	.091	.530
Right Caudate – Right dIPFC	.195	.185	.102	.479
Right Caudate - Left dIPFC	.329	.022	032	.826
Left Caudate – Right vIPFC	086	.559	.223	.120
Left Caudate- Left vIPFC	244	.095	.035	.808
Left Caudate - Right dIPFC	.140	.342	.130	.370
Left Caudate - Left dIPFC	.404	.004*	.089	.539
Right Ventral Striatum – Right vIPFC	.069	.642	306	.031
Right Ventral Striatum - Left vIPFC	.100	.500	0.009	.950
Right Ventral Striatum - Right dIPFC	100	.944	141	.330
Right Ventral Striatum - Left dIPFC	190	.195	141	.330
Left Ventral Striatum - Right vIPFC	.064	.666	348	.012*
Left Ventral Striatum - Left vIPFC	.009	.950	111	.444
Left Ventral Striatum - Right dIPFC	042	.775	049	.737
Left Ventral Striatum- Left dIPFC	221	.130	055	.706

Note: * Survived FDR correction

The full results for the partial correlations, controlling for age, IQ and sex, between CAG repeats and pairwise functional connectivity for both the pre-HD group are displayed in Supplementary Table 4.

Supplementary Table 4. Correlations between all pairwise ROIs and CAG repeats in the pre-HD group.

	Premanifest HD	
	R-Value	P-Value
Right Caudate – Right vIPFC	014	.925
Right Caudate - Left vIPFC	039	.794
Right Caudate – Right dIPFC	.076	.607
Right Caudate - Left dIPFC	.167	.257
Left Caudate – Right vIPFC	032	.828
Left Caudate- Left vIPFC	.013	.930
Left Caudate - Right dIPFC	.104	.483
Left Caudate - Left dIPFC	.166	.260
Right Ventral Striatum – Right vIPFC	033	.823
Right Ventral Striatum - Left vIPFC	.001	.996
Right Ventral Striatum - Right dIPFC	.161	.276
Right Ventral Striatum - Left dIPFC	.031	.833
Left Ventral Striatum - Right vIPFC	.049	.743
Left Ventral Striatum - Left vIPFC	.041	.780
Left Ventral Striatum - Right dIPFC	003	.986
Left Ventral Striatum- Left dIPFC	023	.879

Note: * Survived FDR correction

References

- Scahill, R., Zeun, P., Osborne-Crowley, K., et al. Biological and clinical characteristics of gene carriers far from predicted onset in the Huntington's disease Young Adult Study (HD-YAS): a cross-sectional analysis. *Lancet Neurol* 2020;19:502-512.
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- 3. Penney Jr, J.B., Vonsattel, J.P., Macdonald, M.E., et al. CAG repeat number governs the development rate of pathology in Huntington's disease. *Ann Neurol* 1997;41:689-692.
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- 5. CANTAB® [Cognitive assessment software]. Cambridge Cognition (2019).
- 6. Berg, E.A. A simple objective technique for measuring flexibility in thinking. *J Gen Psychol* 1948;39:15-22.