Behavioral Symptoms in Premanifest Huntington Disease Correlate with Reduced Frontal CB₁R Levels

Jenny Ceccarini¹, Rawaha Ahmad¹, Laura Van de Vliet^{2,3}, Cindy Casteels¹, Mathieu Vandenbulcke^{2,3}, Wim Vandenberghe^{3,4}, and Koen Van Laere¹

¹Nuclear Medicine and Molecular Imaging, University Hospitals Leuven, and Department of Imaging and Pathology, KU Leuven, Leuven, Belgium; ²Department of Old Age Psychiatry, University Psychiatric Centre, KU Leuven, Leuven, Belgium; ³Department of Neurosciences, KU Leuven, Leuven, Belgium; and ⁴Department of Neurology, University Hospitals Leuven, Leuven, Belgium

Many Huntington disease (HD) mutation carriers already have cognitive and psychiatric symptoms in the premanifest (premotor) phase of the disease (pre-HD), but the molecular underpinnings of these symptoms are not well understood. Previous work has shown reduced availability of the cerebral type 1 cannabinoid receptor (CB₁R) in manifest HD. Here, we investigated whether CB₁R binding is related to cognitive and psychiatric symptoms in pre-HD mutation carriers. **Methods:** CB₁R binding was measured with ¹⁸F-MK-9470 (N-[(2S,3S)-3-(3-cyanophenyl)-4-(4-ethoxyphenyl)butan-2-yl]-2methyl-2-(5-methylpyridin-2-yl)oxypropanamide) PET in 15 pre-HD subjects (8 men, 7 women; age, 39.3 ± 9.9 y), 15 gene-negative controls from HD families (9 men. 6 women; age. 37.0 ± 10.6 v). and 12 community controls (6 men and 6 women; age, 39.9 ± 15.1 y). All subjects also underwent extensive assessment of motor and cognitive function, as well as a behavioral test battery including the Problem Behavior Assessment for HD (PBA-HD), and MRI. Parametric binding images of ¹⁸F-MK-9470 were corrected for partial-volume effect. Results: There was no difference in CB1R binding, gray matter volume, cognitive function, or psychiatric scores between gene-negative controls from HD families and community controls, which were therefore pooled to one control group. Compared with controls, pre-HD subjects showed striatal atrophy, a decrease in CB₁R binding in the prefrontal cortex, and higher PBA-HD scores on depression, apathy, and irritability (range, P =0.01-0.005). The PBA-HD scores inversely correlated with CB₁R binding in prefrontal regions and cingulate cortex in pre-HD (range: r = -0.64 to -0.72; P = 0.01-0.008). **Conclusion:** The association between behavioral symptoms and reduced prefrontal CB₁R levels may provide new insight into the molecular basis of neuropsychiatric symptoms in pre-HD and suggest new therapeutic avenues.

Key Words: premanifest HD (pre-HD); cannabinoid type 1 receptor (CB₁R); behavioral symptoms; PET

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Luntington disease (HD) is an autosomal dominant neurodegenerative disorder caused by a CAG repeat expansion in the huntingtin (HTT) gene. The toxic effects of mutant huntingtin

Received Feb. 23, 2018; revision accepted May 24, 2018. For correspondence or reprints contact: Jenny Ceccarini, University Hospital Leuven, Herestraat 49, 3000 Leuven, Belgium.

E-mail: jenny.ceccarini@uzleuven.be

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protein induce neuronal cell loss, most prominently in the caudate nucleus and putamen but also in cortical regions. Clinically, HD patients develop motor dysfunction, cognitive decline, and psychiatric disturbances. Traditionally, the clinical diagnosis of HD is made when a carrier of the HD mutation develops motor signs that are typical of HD. However, many HD mutation carriers already have cognitive changes and psychiatric symptoms, such as depression, apathy, and irritability, in the premotor (premanifest) stage (pre-HD) (1-6). The molecular and anatomic origin of these pre-HD cognitive and psychiatric changes is not well understood.

The type 1 cannabinoid receptor (CB₁R) is a crucial modulator of synaptic transmission in the brain and is involved in the regulation of motor behavior, cognition, and emotion (7). Several lines of evidence suggest an important role for CB₁R in HD. In vivo PET imaging with 18 F-MK-9470 (N-[(2S,3S)-3-(3-cyanophenyl)-4-(4-ethoxyphenyl)butan-2-yl]-2-methyl-2-(5-methylpyridin-2-yl) oxypropanamide) has revealed a substantial reduction of CB₁R binding in patients with manifest HD, even in early manifest stages (8), suggesting that the decline in CB₁R availability may potentially already begin in the pre-HD phase. This hypothesis is supported by the observation that CB₁R protein and messenger RNA levels are decreased in the pre-HD phase in transgenic mouse models of HD, with a further decline during the manifest course of the disease (9). Also, genetic deletion of CB₁R in transgenic HD mouse models aggravates symptomatology, suggesting that the decline of CB₁R in human HD may actually underlie part of the clinical phenomenology.

So far, no human CB₁R PET studies in pre-HD mutation carriers have been reported. In this study, we assessed in vivo CB₁R availability using ¹⁸F-MK-9470 PET in pre-HD mutation carriers in comparison to healthy, age-matched controls. To control for potential effects of the distress caused by growing up in an HD family and undergoing predictive genetic testing for HD (2), we included a group of community controls as well as a control group consisting of gene-negative subjects from HD families.

The first aim of this study was to investigate whether CB₁R availability is already decreased in pre-HD. Second, we wanted to investigate whether CB₁R availability in pre-HD may correlate with specific psychiatric and cognitive symptoms.

MATERIALS AND METHODS

Participants

Fifteen pre-HD subjects (HTT CAG repeat length of 40 or more; absence of chorea or other motor signs typical of HD) were included in the study, as well as 15 gene-negative healthy controls from HD families (CAG repeat length of less than 36) and 12 healthy community

controls (no personal or family history of neuropsychiatric disorders). We included the gene-negative control group in addition to the community control group to control for the distress of growing up in an HD family and undergoing predictive genetic testing for HD. Pre-HD subjects and gene-negative controls were recruited via the Department of Human Genetics of the University Hospitals Leuven after predictive genetic testing for HD. Community controls were recruited through advertisements in community newspapers and on a departmental website.

Subjects were excluded if they met any of the following criteria: past or current abuse of alcohol or other drugs, intake of psychotropic drugs, pregnancy, implanted devices or material that could interfere with the MRI scans, and previous radiation exposure of more than 1 mSv for other medical research studies during the past 12 mo.

All subjects underwent a general physical examination and blood and urine testing to exclude major internal pathologic states. Absence of any drug use at the time of CB₁R PET scanning was confirmed by a urine stick test, which included toxicology for benzodiazepines, neuroleptics, opiates, cocaine and metabolites, amphetamines, and cannabinoids before the CB₁R PET scan in all controls and pre-HD subjects.

The study was approved by the institutional Ethics Committee and was performed in accordance with the latest version of the World Medical Association Declaration of Helsinki. Written informed consent was obtained from all participants before the study.

Clinical Assessments

The motor part of the Unified Huntington's Disease Rating Scale was administered to all subjects on the day of PET. All subjects underwent a neuropsychologic test battery including the Mini-Mental State Examination, a test of episodic memory (Rey Auditory Verbal Learning Test), tests of attention and executive functioning (Digit Span Test and the Digit Symbol Subtest from the Wechsler Adult Intelligence Scale), Trail Making Tests A and B, the Stroop Test, the Letter Verbal Fluency and Animal Verbal Fluency Test, and tests of language (Boston Naming Test) and intelligence (Wechsler Adult Intelligence Scale).

Mood and behavioral symptoms were assessed using the Beck Depression Inventory, the State-Trait Anxiety Inventory, and the Problem Behavior Assessment for HD (PBA-HD) (10,11). The PBA-HD is a semistructured interview specifically designed for assessment of

behavioral problems in HD and comprises 3 subscales: PBA-HD apathy, PBA-HD irritability, and PBA-HD depression. Three of the 15 pre-HD subjects declined to undergo the PBA assessment.

The predicted age of motor onset in the pre-HD group was calculated according to Aylward et al. (12) as $[(-0.81 \times [CAG]_n) + (0.51 \times parental onset age) + 54.87]$. The 10-y probability of disease onset was calculated using the model of Langbehn et al. (13). Disease burden was calculated as $[(CAG]_n - 35.5) \times age]$ (14).

Image Acquisition and Brain Atrophy Assessment

All subjects fasted for at least 4 h before PET imaging. They received 217.4 \pm 68.7 MBq (mean \pm SD) of ¹⁸F-MK-9470 (8,15) as a slow intravenous injection under standard conditions (specific activity, 177.7 ± 86.5 GBq/ μ mol; injected mass dose, 3.7 ± 0.3 μ g). PET imaging was performed in 3-dimensional mode on a HiRez Biograph16 PET/CT camera (Siemens). At the beginning of each PET scan, a low-dose CT scan was conducted for attenuation correction purposes. An ¹⁸F-MK-9470 PET scan of 60 min was started 120 min after injection (16). Images were reconstructed using iterative 3dimensional ordered-subsets expectation maximization and were smoothed afterward with a 3-dimensional isotropic gaussian of 6 mm in full width at half maximum. All subjects also underwent high-resolution MRI on a 3-T scanner (Ingenia; Philips). Individual MR images were used to coregister PET data, to automatically and accurately delineate volumes of interest (VOIs), and to assess potential brain atrophy in patients both at the voxel level (by using voxel-based morphometry) and at the VOI level (by comparing groups' relative gray matter volumes [i.e., gray matter volume normalized to total intracranial volume] according to the 83-VOI N30R83 maximum-probability atlas of Hammers et al. (17)).

Image Processing

Parametric CB_1R binding maps were calculated using the modified SUV [activity concentration \times (subject's body weight + 70/2)/injected activity] (18), a validated noninvasive quantification method that gives a reliable estimate of the total distribution volume of ¹⁸F-MK-9470, as determined by full kinetic modeling in humans (16). The modified SUV measure was chosen for its decreased between-subject variability and therefore increased reproducibility and sensitivity to detect group differences compared with the fractional uptake ratio.

TABLE 1Demographic and Clinical Characteristics of Study Subjects

Characteristic	Community controls	Gene-negative controls	Pre-HD subjects
Subjects	12	15	15
Sex			
Male	6	9	8
Female	6	6	7
Age (y)	39.9 ± 15.1	37.0 ± 10.6	39.3 ± 9.9
[CAG]n	NA	20.33 ± 2.92	42.4 ± 1.8
UHDRS motor	0.7 ± 0.7	1.3 ± 1.1	2.3 ± 2.0
Predicted age of onset (y)	_	_	45.2 ± 4.8
10-y probability of disease onset (%)	_	_	37.1 ± 28.1
Disease burden*			264.3 ± 74.6

^{*}Calculated as ([CAG]n - 35.5) \times age.

NA = not available; UHDRS = Unified HD Rating Scale.

Qualitative data are expressed as n; continuous data are expressed as mean \pm SD.

TABLE 2Neuropsychologic Assessments in Pre-HD Subjects and Controls

Assessment	Control* $(n = 27)$	Pre-HD (n = 15)	Р
Cognition			
MMSE	30.0 (29.0–30.0)	29.0 (28.0–29.0)	0.002
RAVLT total (A1-A5)	55.8 ± 8.6	47.5 ± 10.2	0.00
RAVLT retention (A7)	13.0 (12.0–14.0)	11.0 (7.0–14.0)	0.04
RAVLT recognition (A8)	15.0 (15.0–15.0)	14.0 (12.0–15.0)	0.01
BNT	56.0 (53.0–58.0)	56.0 (53.0–58.0)	NS
TMT A (s)	25.0 (23.0–34.0)	31.0 (18.0–38.0)	NS
TMT B (s)	52.5 (47.5–63.0)	74.0 (54.0–99.0)	0.00
Stroop word reading	43.0 (39.0–45.0)	50.0 (39.0–55.0)	NS
Stroop color naming	50.0 (47.0–60.0)	58.0 (52.0–65.0)	0.03
Stroop interference	82.0 (72.0–91.0)	88.0 (84.0–113.0)	0.02
NART	49.0 (46.0–50.0)	47.0 (42.5–49.0)	NS
DS forward	6.0 (6.0–7.0)	6.0 (5.0–7.0)	NS
DS backward	5.0 (4.0–6.0)	4.0 (4.0–5.0)	0.0
VF animal	22.0 (21.0–26.0)	19.0 (16.0–27.0)	NS
VF N	13.0 (10.0–15.0)	9.0 (5.0–10.0)	0.0
VF A	13.0 (11.0–15.0)	8.0 (6.0–10.0)	0.00
VF K	14.33 ± 4.64	11.53 ± 6.31	NS
Digit symbol	85.0 (76.0–93.0)	64.0 (54.0–88.0)	NS
Block patrons	67.0 (65.0–68.0)	66.0 (57.0–67.0)	NS
lood and behavior			
BDI	1.0 (0.0–5.0)	4.0 (1.0–10.0)	NS
STAI 1	47.0 (44.0–55.0)	49.0 (43.0–50.0)	NS
STAI 2	47.8 ± 3.9	46.5 ± 6.1	NS
PBA-HD total	2.7 ± 3.1	10.0 ± 7.7	0.00
PBA-HD apathy	0.1 ± 0.3	1.4 ± 2.0	0.00
PBA-HD irritability	0.4 ± 0.6	1.5 ± 1.2	0.0
PBA-HD depression	0.9 ± 1.3	3.2 ± 3.0	0.02

^{*}Gene-negative and community controls.

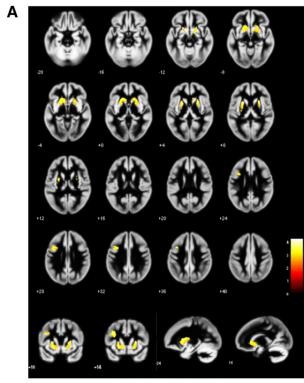
MMSE = Mini-Mental State Examination; RAVLT = Rey Auditory Verbal Learning Test; BNT = Boston Naming Test; NS = not significant; TMT = Trail Making Test; NART = National Adult Reading Test; DS = Digit Span Test; VF N, A, K = verbal fluency for letters N, A, and K; BDI = Beck Depression Inventory; STAI = State-Trait Anxiety Inventory.

Skewed data are median followed by quartiles 1–3 in parentheses. Normally distributed data are mean \pm SD. P values are for results surviving Kruskal–Wallis test and Bonferroni test.

Further image processing was performed using an automatic pipeline in the PMOD brain PNEURO tool (PMOD, version 3.6; PMOD Technologies) (19). PET images were first rigidly coregistered to the corresponding volumetric T1-weighted MR images. The individual T1-weighted MR images were then nonlinearly coregistered to the standard Montreal Neurologic Institute space MRI template. Subsequently, the same was done for the PET images using the same spatial transformation as for the registered MR images. T1-weighted MR images were segmented into gray matter, white matter, and cerebrospinal fluid within native MRI space to generate a total of 83 individual VOIs according to the N30R83 atlas (17). Automatic delineation of the deep nuclei was performed by T1-weighted MRI parcellation in the PMOD PNEURO tool. Partial-volume effect (PVE) correction was performed as implemented in PNEURO (20), using the methods of Müller-Gärtner et al. (21) and Rousset et al. (22) for the voxelwise and VOI-based analysis, respectively.

Image Analysis and Statistics

First, a voxelwise analysis was performed using Statistical Parametric Mapping (SPM8; Wellcome Department of Cognitive Neurology). PVE-corrected 18 F-MK-9470 parametric maps were smoothed with a 10-mm isotropic gaussian kernel and were masked to exclude extracerebral activity. For analysis of absolute 18 F-MK-9470 binding, no global normalization was used. t maps were interrogated at a cluster-level P value (P_{cluster}) of less than 0.05 and a voxel-level P value (P_{height}) of less than 0.001 and with an extent threshold (k_{ext}) of 50 voxels. A VOI-based group-comparison analysis was also performed using the N30R83 atlas. To avoid type II errors, VOIs were grouped into composite, bilateral VOIs for the frontal cortex, parietal cortex, lateral temporal cortex, medial temporal cortex, occipital cortex, anterior cingulate cortex, posterior cingulate cortex, and striatum. Because of its potential importance for the cognitive and behavioral



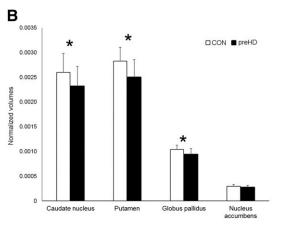


FIGURE 1. Decrease in striatal gray matter volume in pre-HD subjects in comparison with healthy controls (CON) revealed at voxel level ($P_{\rm height} < 0.005$) (A) and at VOI level (B). *P < 0.05.

measures in this study, the frontal cortex was also subdivided into prefrontal, dorsolateral prefrontal, ventrolateral prefrontal, ventromedial prefrontal, and orbitofrontal.

Statistical analyses were performed using Statistica (version 12; Statsoft Inc.) and SPSS (version 22.0; IBM). Shapiro–Wilk tests were used to test the distribution of variables against normality. ANOVA or the Kruskal–Wallis test, followed by a Bonferroni post hoc test, was used as appropriate.

To study the interrelatedness of the different behavioral and cognitive measures, we performed a factor analysis (principal components for the method extraction, 5 factors extracted, varimax normalized for the factor rotation).

Moreover, we investigated the degree to which regional CB₁R availability in the pre-HD group was associated with cognitive and psychiatric measures when significantly different from controls, performing an

SPM correlation analysis ($P_{\rm cluster}$ < 0.05 and at a voxel-level $P_{\rm height}$ < 0.001, unless otherwise stated, with a $k_{\rm ext}$ of 50 voxels) and a VOI-based analysis (Spearman rank test, P < 0.05).

RESULTS

Demographic and Clinical Characteristics

In total, 15 pre-HD subjects, 15 gene-negative controls, and 12 community controls were included in the study (Table 1). Unified Huntington's Disease Rating Scale motor scores were slightly higher (P = 0.032) in pre-HD subjects than in community controls (Table 1), consistent with the concept that subtle motor changes can be detected on a group level in the pre-HD phase (5). There was no difference in age among the 3 groups (range, P > 0.64). Moreover, no statistically significant difference in sex distribution was observed among the 3 groups (χ^2 test, P = 0.86).

There were no differences between community controls and gene-negative controls in neuropsychologic scores, gray matter volume, or CB₁R binding (Supplemental Table 1; Supplemental Figs. 1 and 2 [supplemental materials are available at http://jnm.snmjournals.org]). The community controls and gene-negative controls were therefore pooled into one control group for comparison with pre-HD subjects (Table 2).

Regarding the cognitive assessment, pre-HD subjects performed worse than controls on MMSE (P = 0.002), Rey Auditory Verbal

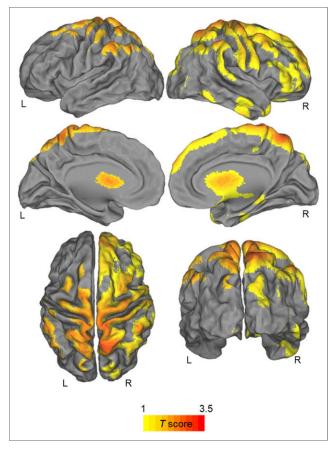


FIGURE 2. Statistical parametric t map of significantly decreased PVE-corrected CB₁R availability in pre-HD mutation carriers vs. controls. Maps are projected on PALS-B12 atlas (a population-averaged landmark- and surface-based atlas derived from 12 healthy young adults). Significance is shown with t statistic color scale corresponding to level of significance at voxel level.

Learning Test (range, P=0.008-0.04), verbal fluency (range, P=0.007-0.03), Stroop color naming (P=0.03), and Stroop interference (P=0.02) (Table 2). As for mood and behavioral measures, pre-HD subjects had a significantly higher PBA-HD total score than controls (P=0.002), as well as higher subscores (PBA-HD apathy [P=0.005], PBA-HD irritability [P=0.01], and PBA-HD depression [P=0.02]) (Table 2). All reported P values survived Bonferroni post hoc correction.

Gray Matter Volume

Voxel-based morphometry revealed significant striatal gray matter atrophy in pre-HD subjects bilaterally in the putamen, caudate nucleus, nucleus accumbens, and left middle frontal gyrus (Fig. 1A; Supplemental Table 2). These volumetric gray matter reductions in the pre-HD group were also observed at the VOI level by comparing relative VOIs (caudate: -10.8%, P=0.03; putamen: -11.5%, P=0.002; pallidum: -9.5%, P=0.005; nucleus accumbens: -7.6%, P=0.065; Fig. 1B). In pre-HD subjects, there were no significant associations between either the normalized volume decrease or the percentage decrease in gray matter volume and the cognitive and behavioral measures (nonparametric Spearman correlation test, P<0.05).

CB₁R Availability and Clinical Characteristics in Pre-HD

VOI-based analysis showed no significant CB_1R differences between the 2 control groups (gene-negative controls vs. community controls) in any region (average difference, $-1.0\% \pm 1.8\%$; range, P=0.5–0.9) (Supplemental Fig. 2). Therefore, also here the community control group was pooled with the gene-negative control group to create one control group for comparison of the 18 F-MK-9470 data with the pre-HD group.

Compared with controls, the pre-HD group showed a decrease in PVE-corrected CB₁R availability in the superior and middle prefrontal gyrus, superior parietal gyrus, precentral and posterior gyrus ($P_{\text{cluster}} = 0.003$ and $P_{\text{height}} = 0.001$ for voxel level, $k_{\text{ext}} = 14449$), and caudate nucleus ($P_{\text{cluster}} = 0.016$ and $P_{\text{height}} = 0.001$ for voxel level, $k_{\text{ext}} = 205$) (Fig. 2). Similar results were obtained when the voxel-based analysis was performed with nonsmoothed PVE-corrected modified SUV images.

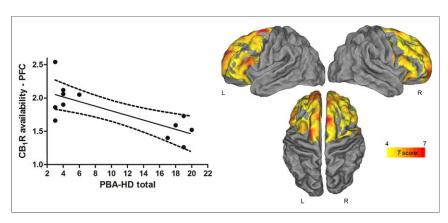


FIGURE 3. Negative correlation between CB₁R and PBA-HD total in pre-HD. (Left) Scatterplot showing inverse correlation between individual PBA-HD total scores and CB₁R availability in prefrontal cortex (PFC) at VOI-based level. Solid line represents regression line, and dotted lines represent 95% confidence interval. (Right) Statistical parametric maps showing inverse correlation between PVE-corrected CB₁R availability in pre-HD and PBA-HD total. Maps are projected on PALS-B12 atlas (a population-averaged landmark- and surface-based atlas derived from 12 healthy young adults). Significance is shown with *t* statistic color scale corresponding to level of significance at voxel level.

These findings were largely confirmed at a VOI level. Total PBA-HD scores showed an inverse correlation with CB₁R availability in all prefrontal regions (range, r = -0.64 to -0.72; range, P = 0.02–0.008; Fig. 3, left) and cingulate cortex (posterior cingulate cortex: r = -0.60, P = 0.04; anterior cingulate cortex: r = -0.69, P = 0.01) of pre-HD subjects. Regarding the PBA-HD subscores, only the correlation between the PBA-HD depression score and CB₁R availability reached significance in the prefrontal cortex (r = -0.61, P = 0.03; Fig. 4, left), orbitofrontal cortex, and anterior cingulate cortex (r = -0.59, P = 0.04) (PBA-HD apathy: range, P = 0.09–0.21; PBA-HD irritability: range, P = 0.26–0.59).

Significant factor loadings were found for the Beck Depression Inventory score, State-Trait Anxiety Inventory 2, Total PBA-HD, and PBA-HD apathy and depression (range of marked loadings, 0.74–0.88), showing that mood and motivation symptoms correlated with each other in the pre-HD group. Individual scores for factor 1 were intercorrelated with the CB₁R availability in all prefrontal cortex regions (range, r = 0.60–0.69; P = 0.013–0.41) and anterior cingulate cortex (r = 0.67, P = 0.018) (Supplemental Fig. 3).

 ${\rm CB_1R}$ availability did not correlate with [CAG]n (SPM correlation analysis with [CAG]n as covariate and age as nuisance variable), even at a liberal $P_{\rm height}$ threshold of less than 0.01 (uncorrected). There were no significant correlations between disease

burden and either regional CB₁R levels (range, P = 0.27–0.97) or PBA-HD scores (range, P = 0.25–0.75) in the pre-HD group.

DISCUSSION

Psychiatric symptoms are common in pre-HD (*1*–6), but the underlying brain changes are poorly understood. To our knowledge, our PET study is the first to find a molecular correlate of behavioral symptoms in pre-HD.

In contrast to the more global decrease in cerebral CB₁R availability seen in manifest HD patients (8), we found regionally restricted CB₁R reductions in our pre-HD group in the superior and middle prefrontal gyrus, precentral gyrus, and caudate nucleus. These regional CB₁R reductions in pre-HD are unlikely to be caused by the psychologic stress associated with growing up in an HD family and undergoing predictive genetic testing, because there was

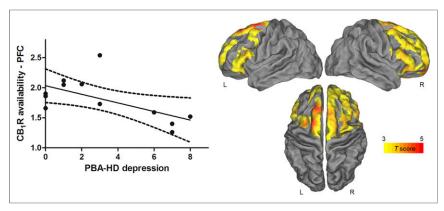


FIGURE 4. Negative correlation between CB_1R and PBA-HD depression in pre-HD. (Left) Scatterplot shows inverse correlation between individual PBA-HD depression scores and CB_1R availability in prefrontal cortex (PFC) at VOI-based level. Solid line represents regression line, and dotted lines represent 95% confidence interval. (Right) Statistical parametric maps and inverse correlation between PVE-corrected CB_1R availability in pre-HD mutation carriers with PBA-HD depression. Maps are projected on PALS-B12 atlas (a population-averaged landmark- and surface-based atlas derived from 12 healthy young adults). Significance is shown with t statistic color scale corresponding to level of significance at voxel level.

no difference in CB_1R availability between gene-negative controls and community controls. No differences were found in CB_1R availability levels, cognitive measures, or psychiatric symptoms between the 2 control groups, indicating that growing up with affected and at-risk family members and undergoing predictive genetic testing for HD did not have any detectable influence on CB_1R signaling and cognitive and behavioral measures. Instead, the CB_1R reductions in pre-HD may be due to a direct repressive effect of mutant huntingtin on CB_1R gene transcription (23,24).

Importantly, we found CB₁R in the prefrontal and cingulate cortex to have an inverse correlation with behavioral symptoms, in particular depression. The prefrontal cortex and anterior cingulate cortex are involved in emotion regulation via reciprocal connections with limbic structures and play a major role in the pathophysiology of mood disorders (25,26). Moreover, a large body of preclinical and human data suggest an important role for hypofunction of the endocannabinoid system, particularly in the prefrontal cortex, in the pathogenesis of depression (27,28). Genetic deletion or pharmacologic inhibition of CB₁R in rodents induces symptoms that mimic human depression (27,29). Conversely, activation of endocannabinoid transmission has antidepressantlike properties in established animal models of depression (30,31). Finally, clinical trials showed that the CB₁R inverse agonist rimonabant increases the risk of depressed mood (32,33).

Although we did not find a significant correlation between PBA-HD apathy and CB₁R availability, there was a strong intercorrelation between test scores for depression and apathy using factor analysis, suggesting that apathy and depression are difficult to tease apart in pre-HD and that the neurodegenerative process disrupts circuits involved in both mood and motivation at an early stage.

If the frontal CB₁R reduction causally contributes to the behavioral symptoms in HD mutation carriers, pharmacologic endocannabinoid system activation could have beneficial effects on these symptoms. Interestingly, a pilot study with the CB₁R agonist nabilone in HD patients found preliminary evidence for improvement of neuropsychiatric symptoms (including depression and apathy) (34).

Some limitations of the present study should be acknowledged. First, the pre-HD group was relatively small and needs independent replications in larger pre-HD populations. Second, longitudinal follow-up studies in the pre-HD cohort will be required to assess the potential value of CB₁R PET as a biomarker for pre-HD progression in comparison with other imaging modalities such as volumetric MRI (*35*) and PDE10A PET (*36–38*). Finally, larger clinical studies are warranted to investigate the effects of activators of endocannabinoid signaling on behavioral symptoms in pre-HD and HD.

CONCLUSION

The principal findings of this PET study are the decreased CB_1R availability in the prefrontal cortex in pre-HD mutation carriers and the strong inverse correlation between frontal CB_1R levels in pre-HD and

behavioral symptoms, in particular scores on depression. Taken together, the association between behavioral symptoms and reduced prefrontal CB₁R levels may provide new insight into the molecular basis of neuropsychiatric symptoms in pre-HD and suggest new therapeutic avenues targeting the endocannabinoid system.

DISCLOSURE

Jenny Ceccarini is a postdoctoral fellow of the Research Fund–Flanders. Wim Vandenberghe and Koen Van Laere are Senior Clinical Investigators of the Research Fund–Flanders. No other potential conflict of interest relevant to this article was reported.

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