

Review

The Role of Hypothalamic Pathology for Non-Motor Features of Huntington's Disease

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Abstract. Huntington's disease (HD) is a fatal genetic neurodegenerative disorder. It has mainly been considered a movement disorder with cognitive symptoms and these features have been associated with pathology of the striatum and cerebral cortex. Importantly, individuals with the mutant huntingtin gene suffer from a spectrum of non-motor features often decades before the motor disorder manifests. These symptoms and signs include a range of psychiatric symptoms, sleep problems and metabolic changes with weight loss particularly in later stages. A higher body mass index at diagnosis is associated with slower disease progression. The common psychiatric symptom of apathy progresses with the disease. The fact that non-motor features are present early in the disease and that they show an association to disease progression suggest that unravelling the underlying neurobiological mechanisms may uncover novel targets for early disease intervention and better symptomatic treatment. The hypothalamus and the limbic system are important brain regions that regulate emotion, social cognition, sleep and metabolism. A number of studies using neuroimaging, postmortem human tissue and genetic manipulation in animal models of the disease has collectively shown that the hypothalamus and the limbic system are affected in HD. These findings include the loss of neuropeptide-expressing neurons such as orexin (hypocretin), oxytocin, vasopressin, somatostatin and VIP, and increased levels of SIRT1 in distinct nuclei of the hypothalamus. This review provides a summary of the results obtained so far and highlights the potential importance of these changes for the understanding of non-motor features in HD.

Keywords: Huntington's disease, hypothalamus, orexin, hypocretin, oxytocin, vasopressin

NON-MOTOR FEATURES OF HUNTINGTON'S DISEASE

Huntington's disease (HD) is more than a movement disorder. It is always caused by an expanded CAG repeat in the huntingtin (*HTT*) gene. A positive gene test in combination with unequivocal signs of motor disturbances are required for the clinical diagnosis of HD today (i.e., manifest HD) [1]. However,

a large number of clinical studies have demonstrated the presence of non-motor features decades before the manifestation of the movement disorder and alterations using neuroimaging techniques in individuals who carry the mutant *HTT* gene (i.e., premanifest HD [1–4]). The motor disturbances in HD have been associated with progressive and pronounced dysfunction and pathology in the striatum of the basal ganglia and the cerebral cortex, the most affected areas in HD. Cognitive changes are common and constitute deficits in executive function with reduced capacity for concentration, attention, mental flexibility and organization. These cognitive changes have also been

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associated with pathology in the striatum and the cerebral cortex. Other non-motor features of HD include a wide range of psychiatric symptoms such as apathy, depression, anxiety, irritability and altered social cognition [5, 6]. Reduced recognition of facial expression of emotions has also been detected in pre-manifest HD [7–10]. Reduced emotional recognition has recently been found to be associated with apathy, which is the psychiatric symptom that most clearly increases with disease progression [11, 12]. Sleep problems and an altered circadian rhythm occur in many individuals with HD [13–18]. Metabolic alterations include increased appetite with a higher caloric intake being necessary to counterbalance weight loss in HD [19, 20]. A high body mass index (BMI) at motor onset has been associated with a slower disease progression [21]. The underlying neurobiological mechanisms for these non-motor features are not known. As their occurrence is common early in the disease process and have associations to disease progression, further understanding of how they arise and develop may lead to important insight into early disease mechanisms and uncover new targets for disease modification.

The regulation of emotion, sleep and metabolism is governed by the hypothalamus and the limbic system. The hypothalamus is made up of interconnected nuclei that receive inputs both from the periphery, e.g., thyroid hormones, leptin, ghrelin and insulin, and the central nervous system (CNS). Besides regulating the endocrine axes of the body, its many different neuropeptide-expressing neuronal populations project within the region and to other areas of the brain to regulate emotion, sleep and metabolism. The hypothalamus is part of the larger limbic system that includes the hippocampus, gyrus cinguli, pre-frontal cortex, insula, septal nuclei, amygdala, ventral striatum, ventral tegmental area and raphe nucleus [22, 23]. In light of the presence of non-motor features in HD, this system has gained increasing interest for investigations of changes in clinical material and experimental models of HD (previously reviewed in [24–27]). These authors reviewed the state of knowledge of this area in HD in 2012 in this journal [28] and provides here an updated overview of the major findings made in the hypothalamus and the limbic system in HD. The review is based on a literature search on the PubMed database up to 2019 with the search terms ‘Huntington disease’, ‘huntingtin’, ‘hypothalamus’, ‘limbic system’, ‘orexin’, ‘hypocretin’, ‘oxytocin’, and ‘vasopressin’.

HYPOTHALAMIC CHANGES IN CLINICAL HD DETECTED BY NEUROIMAGING STUDIES

A combination of neuroimaging studies and post-mortem analyses of human hypothalamic tissue have identified alterations in the hypothalamic region in HD (Fig. 1, Table 1). Studies of the hypothalamic region in general are challenged by practical factors such as the scarcity of such tissue in brain banks and by intrinsic factors such as the difficulties to define the whole structure or specific nuclei within the hypothalamus due to the lack of clear anatomical borders. Nevertheless, structural analyses using voxel-based morphometry and mathematical modelling based on grey matter signals in the hypothalamic region have detected significant differences between premanifest HD and age- and sex-matched controls in several studies [18, 29–31]. Studies using positron emission tomography (PET) have found increased microglia activation and reductions in dopamine D2 receptors also in premanifest HD [32, 33]. Attempts to estimate the hypothalamic volume have not revealed any significant differences between HD and controls [34]. Nevertheless, these studies indicate that the hypothalamic region is affected early in clinical HD.

NEUROPATHOLOGICAL CHANGES IN THE HYPOTHALAMUS IN CLINICAL HD

Postmortem analyses of hypothalami from individuals with HD have revealed a number of changes in different nuclei of this region (Fig. 1, Table 1). The hypothalamus constitutes several nuclei such as the paraventricular nucleus (PVN), supraoptic nucleus (SON), infundibular nucleus, ventromedial nucleus of the hypothalamus (VMH), suprachiasmatic nucleus (SCN), nucleus tuberalis lateralis (NTL), lateral hypothalamic area (LHA) and mammillary bodies. The frequency of huntingtin inclusions as assessed using the EM48 antibody has been estimated to be between 0.5–2.8% in the different nuclei [35]. Although stereological estimation for the total number of neurons and cells in the whole hypothalamus in sections stained for the Nissl stain cresyl violet did not reveal any significant differences between HD and control cases, a specific loss of neurons has been detected in the PVN by 23% and in the NTL by 32% in HD tissue [35, 36]. Early analyses have also indicated a reduced number

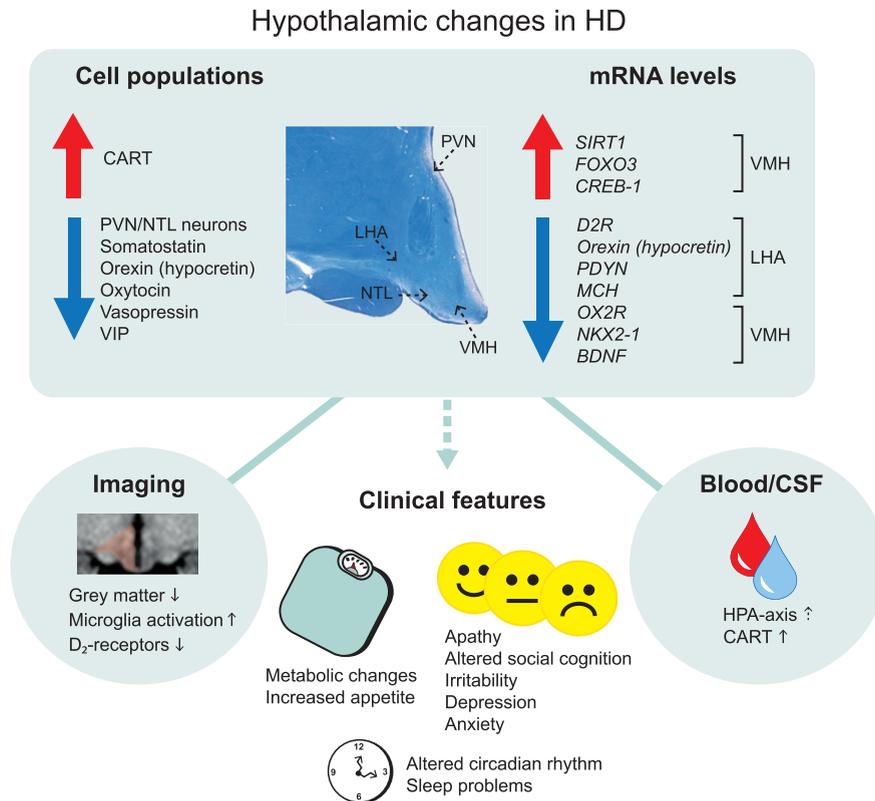


Fig. 1. Overview of main hypothalamic changes in Huntington's disease. Altered immunoreactivity and gene expression levels of cell populations detected in specific hypothalamic nuclei. Changes that result in upregulation and downregulation are indicated by the arrows. The location of specific hypothalamic nuclei is indicated by the dashed arrows on the cross-sectional human hypothalamic section stained for Cresyl violet and Luxol fast blue. Hypothalamic changes are also detected in the blood and cerebrospinal fluid as well as from different imaging paradigms. These changes are thought to be a key contributor to the clinical features in Huntington's disease. CART, cocaine and amphetamine regulated transcript; CREB-1, cyclic AMP-responsive element-binding protein 1; D2R, dopamine D2 receptor; FOXO3, Forkhead box O3; HPA, hypothalamic–pituitary–adrenal; LHA, lateral hypothalamic area; MCH, melanin-concentrating hormone; NTL, nucleus tuberalis lateralis; NKX2-1, NK2 homeobox 1; OX2R, Orexin 2 receptor; PVN, paraventricular nucleus; PDYN, prodynorphin, SIRT1, sirtuin 1; VIP, vasoactive intestinal peptide; VMH, ventromedial hypothalamus.

of somatostatin neurons in the NTL [37–39]. The function of this nucleus is still not well understood. Another study found a reduced number of neurons expressing oxytocin by 45% and vasopressin by 24% in HD cases compared to controls [35]. The number of oxytocin and vasopressin immunopositive neurons were reduced in a case of premanifest HD with Vonsattel grade 0, suggesting that loss of oxytocin and vasopressin may occur early in HD [40]. Oxytocin has been implicated in the ability to recognize facial expression of emotion, which has been found to be reduced already in premanifest HD [41, 42]. Acute intranasal administration of oxytocin to HD gene carriers normalized their altered brain activation pattern in response to recognizing facial

expression of emotional disgust, indicating a role for oxytocin in altered emotion recognition in HD [43]. As for vasopressin, it is not known whether vasopressin levels or their activity are altered in HD. The vasopressin system has been linked to aggression in animals where aggressive behavior can be inhibited by vasopressin V1a receptor antagonists [44, 45]. There is an ongoing Phase 2 clinical trial with Azevan Pharmaceuticals using a vasopressin receptor antagonist (SRX246) against irritability in manifest HD (<https://clinicaltrials.gov/ct2/show/NCT02507284>).

Several studies have investigated the LHA in HD. The LHA controls several functions that are disturbed in HD and pathology in this area could therefore be a contributing if not causative factor for part

Table 1
Summary of main findings in the hypothalamus in clinical HD

Pathology	HD stage	References
Atrophy	↑ Premanifest and manifest (VBM)	[29, 31, 121]
Grey matter volume	↓ Premanifest and manifest (VBM)	[30, 31, 122]
	= Premanifest and manifest	[34]
White matter volume	↓ Premanifest and manifest (DTI)	[122, 123]
Fornix demyelination	↑ Premanifest and manifest (DTI, VBM)	[124]
NTL volume/neurons	↓ Cell counts and volume (HD grade 2–4)	[35–38]
PVN neurons	↓ Cell counts (HD grade 2–4)	[35]
Microglial activation	↑ Premanifest and manifest (PET)	[32, 33]
Functioning of HPA axis	↑ Premanifest and manifest	[107, 109, 125, 126]
Circadian rhythm dysfunction	↑ Premanifest and manifest	[127]
AgRP	= Immunoreactivity (HD grade 1–4)	[128]
BDNF	↓ mRNA levels (HD grade 1–4)	[52]
CART	↑ Immunoreactivity (HD grade 2–4)	[35]
	= Immunoreactivity, mRNA levels (HD grade 0, HD grade 1–4)	[40, 52, 128, 129]
CREB-1	↑ mRNA (HD grade 1–4)	[52]
CRH	= Immunoreactivity (HD grade 1–4)	[129]
	↑ mRNA (HD grade 1–4)	[129]
Dopamine D2 receptor	↓ Premanifest and manifest (PET), mRNA levels (HD grade 1–4)	[32, 52]
FOXO3	↑ mRNA (HD grade 1–4)	[52]
MCH	= Immunoreactivity (HD grade 2–4)	[51, 52]
	↓ mRNA levels (HD grade 1–4)	[52]
NPY	= Immunoreactivity (HD grade 2–4)	[35]
	↓ Immunoreactivity (HD grade 1–4)	[128]
	↑ mRNA (HD grade 1–4)	[128]
Orexin	= Immunoreactivity (HD grade 0)	[40]
	↓ Immunoreactivity, mRNA levels (HD grade 1–4)	[35, 51, 52, 130]
Orexin 2 receptor	↓ mRNA levels (HD grade 1–4)	[52]
Oxytocin	↓ Immunoreactivity (HD grade 0)	[40]
	↓ Immunoreactivity (HD grade 2–4)	[35]
	= Immunoreactivity (HD grade 1–4)	[129]
Prodynorphin	↓ mRNA levels (HD grade 1–4)	[52]
SIRT1	↑ mRNA (HD grade 1–4)	[52]
Somatostatin	↓ Immunoreactivity (mid-late HD)	[39]
TRH	= mRNA levels (HD grade 1–4)	[52, 129]
Tyrosine hydroxylase	↓ mRNA levels (HD grade 1–4)	[52]
Vasopressin	↓ Immunoreactivity (HD grade 0)	[40]
	↓ Immunoreactivity (HD grade 1–4)	[35, 57]
	= Immunoreactivity (HD grade 1–4)	[129]
	= mRNA levels (HD grade 1–4)	[57]
VIP	↓ Immunoreactivity (HD grade 1–4)	[57]
	= mRNA (HD grade 1–4)	[57]

AgRP, agouti-related protein; BDNF, brain-derived neurotrophic factor; CART, cocaine and amphetamine regulated transcript; CREB-1, cyclic AMP-responsive element-binding protein 1; CRH, corticotrophin releasing hormone; DTI, diffusion tensor imaging; FOXO3, forkhead box O3; MCH, melanin-concentrating hormone; NPY, neuropeptide Y; PET, positron emission tomography; SIRT1, sirtuin 1; TRH, thyrotropin-releasing hormone; VBM, voxel based morphometry; VIP, vasoactive intestinal peptide.

of the clinical manifestation of the disease. Experimental lesions of the LHA established the “lateral hypothalamic syndrome” characterized by hypophagia, adipisia, hypoactivity and weight loss [46–48]. Later studies established an important role for the LHA in the regulation of sleep, energy balance, reward and motivated behaviors [49]. In HD, there is a 38% reduction in the number of orexin (hypocretin) immunopositive-neurons in HD and mRNA levels are also diminished compared to control cases [35, 50–52]. Orexin is a neuropeptide that is only

expressed in the LHA and it plays an important role in the regulation of sleep, emotion and metabolism [53, 54]. Melanin-concentrating hormone (MCH) is another emotion-regulating neuropeptide that is expressed in neighboring neurons within the LHA [55]. A recent study showed significant reduction of MCH mRNA levels in HD cases compared to controls although the number of MCH-immunopositive neurons is not affected [52]. Analyses of mRNA levels of other factors in the LHA in HD cases indicated that the dopamine D2-receptor (D2R) levels are also

reduced [52]. mRNA levels of D2R are not reduced in other hypothalamic areas suggesting that the signal of D2R reductions detected using PET may arise from the LHA specifically.

Sleep problems and altered circadian rhythm are common in patients with HD [13–18, 56]. The SCN is the key brain region for regulating sleep and circadian rhythm. Postmortem analyses of the SCN from patients with HD have shown that numbers of vasoactive intestinal peptide (VIP)- and vasopressin-immunoreactive neurons were reduced by 85% and 33% respectively in HD cases compared to controls [57]. This occurred in the presence of normal mRNA levels of these genes as well as unaffected number of neurons expressing melatonin receptors in this area [57]. Hence, there is specific pathology in the SCN in HD that may contribute to alterations in sleep and circadian rhythm.

HYPOTHALAMIC GENE EXPRESSION CHANGES IN CLINICAL HD

Alterations in specific hypothalamic genes have also been reported in HD (Fig. 1, Table 1). A recent study showed reduced expression of mRNA levels of brain-derived neurotrophic factor (*BDNF*) in the VMH but not in the LHA, PVN or SON of HD cases compared to controls [52]. The VMH constitutes a group of cells involved in the regulation of energy homeostasis as well as aggression [58–64]. *BDNF* is expressed in the VMH [65] and reduced expression of *BDNF* in mice has been associated with aggressiveness, hyperphagia and obesity [66, 67]. *BDNF* is well known to be reduced in the cerebral cortex in clinical HD and has been implicated in the pathogenesis in HD [52, 68]. Hence, the effects of the *BDNF* system specifically in the VMH may also play a role in HD.

The metabolic regulators and energy sensors sirtuins (*SIRT*) have been implicated in HD. Neuroprotective effects of *SIRT1* inhibitors have been shown in drosophila, mammalian cells and mouse models of HD [69]. One initial clinical trial to evaluate safety has been conducted using the *SIRT1* inhibitor, selisistat, in HD patients [70]. *SIRT1* acts in the hypothalamus to control metabolism and longevity partly through the regulation of maturation of hypothalamic peptide hormones [71–74]. A recent study showed an increase in mRNA levels and immunoreactivity of *SIRT1* in both the LHA and VMH apart from other affected brain regions in

HD such as the striatum and cerebral cortex [52]. No changes were found in the less affected cerebellum. Analyses of mRNA levels of downstream targets of *SIRT1* such as *FOXO3* indicated increased levels in the LHA and VMH in HD. *FOXO3* has also been shown to be upregulated in the striatum in HD [75, 76]. This data indicate that the *SIRT1* system is affected in sensitive brain regions in HD and that the hypothalamic effects of these factors need to be considered in future studies targeting this system in HD.

OTHER LIMBIC SYSTEM CHANGES IN CLINICAL HD

Although it is beyond the scope of this review to discuss in detail the alterations of the limbic system in HD, it is important to consider the interactions with other limbic structures which form widely distributed networks for a proper perspective. While the definition of the limbic system is not always clear, the anatomical structures generally accepted today and included in this review comprises of the hippocampus, amygdala, ventral striatum, nucleus accumbens, raphe nucleus, ventral tegmental area, cingulate cortex and prefrontal cortex [22, 23]. Table 2 summarizes the main alterations detected to date in the limbic system in clinical HD and in rodent models. Imaging studies using voxel based morphometry (VBM) in magnetic resonance images (MRI) have indicated increased atrophy and grey matter loss in these limbic structures already in premanifest HD [77–86]. Whereas histopathological studies on the limbic structures in HD are scarce, there is evidence for cell loss in the dorsal raphe nucleus, anterior cingulate cortex and prefrontal cortex in symptomatic HD cases with Vonsattel grade 1–4 [83, 87, 88]. Functional MRI (fMRI) studies have reinforced our understanding of limbic system dysfunction in HD. Reduced functional connectivity, network integrity and activity was found in the HD hippocampus, amygdala, ventral striatum, cingulate cortex and prefrontal cortex using different fMRI paradigms assessing verbal working memory, emotional processing, interference/conflict resolution and attention/alertness [89–99]. Such alterations in activity patterns produce multidimensional maps that to some extent reflect internal states of brain processing in response to a task. These changes are already present in the premanifest stage suggesting that early changes in the limbic structures might contribute in part to disease pathogenesis. As cognition and

Table 2
Summary of main findings in the limbic system in clinical HD and rodent HD models

Region/pathology	HD stage	HD rodent models	References
<i>Hippocampus</i>			
Atrophy	↑ Premanifest	Not assessed	[81]
Network integrity	↓ Manifest	Not assessed	[99]
Volume	= Premanifest	= VBM (R6/2)	[82, 131, 132]
	↓ Premanifest and manifest		[78–80]
Cerebral blood flow	↑ Premanifest	Not assessed	[91]
Neurogenesis	Not assessed	↓ R6/2, R6/1, N171-82Q, YAC128	[133–138]
Androgen receptor	Not assessed	↓ Protein levels (R6/1)	[139]
BDNF	Not assessed	↓ mRNA and protein levels (R6/1), Hdh ^{Q111/Q111} , zQ175	[140–143]
Dendritic spine density	Not assessed	↓ R6/2	[144]
Ferritin	Not assessed	↑ Immunoreactive neurons (R6/2)	[145]
PSA-NCAM	Not assessed	↓ R6/1, R6/2	[146]
Serotonin	Not assessed	↓ R6/2	[147]
Serotonin receptors	Not assessed	↓ R6/1	[148]
<i>Amygdala</i>			
Atrophy	↑ Premanifest	tgHDrat	[81, 149]
Volume	= Premanifest	= VBM (R6/2)	[82, 132]
	↓ Premanifest and manifest		[77]
Functional connectivity	↓ Premanifest	Not assessed	[98]
Dopamine D2 receptor	↓ Manifest	Not assessed	[150]
TRH	↑ Peptide levels (manifest)	Not assessed	[151]
<i>Ventral striatum/nucleus accumbens</i>			
Volume	↓ Premanifest and manifest (HD stage 1-2)	Not assessed	[78, 80, 82, 152]
Functional connectivity	Not assessed	↓ BACHD rats	[153]
Activity	↓ fMRI (premanifest)	Not assessed	[96, 97]
BDNF	Not assessed	↓ mRNA levels (BACHD rats)	[154]
Serotonin	Not assessed	↓ Immunoreactivity (tgHD rat)	[155]
Somatostatin	↑ Peptide levels, immunoreactivity (manifest)	Not assessed	[151, 156]
Tyrosine hydroxylase	Not assessed	↑ Immunoreactivity (tgHD rat)	[155]
<i>Raphe nucleus</i>			
Volume	↓ Manifest	Not assessed	[83]
Neuronal counts	↓ Manifest	Not assessed	[83]
Serotonin	Not assessed	↓ Immunoreactivity (tgHD rat)	[155]
Tyrosine hydroxylase	Not assessed	↑ Immunoreactivity (tgHD rat)	[155]
<i>Ventral tegmental area</i>			
Dopaminergic cell fate	Not assessed	↑ Gene expression (tgHD rat)	[157]
Tyrosine hydroxylase	Not assessed	↑ Immunoreactivity (tgHD rat)	[158]
<i>Cingulate cortex</i>			
Volume	↓ Premanifest and manifest (VBM)	↓ R6/1	[80, 84, 85, 159]
Functional connectivity	↓ Premanifest and manifest (fMRI)	Not assessed	[92]
Activity	↓ Premanifest and manifest (fMRI)	Not assessed	[93–95, 160, 161]
	↑ Premanifest and manifest (fMRI)		[162–164]
Neuronal count	↓ Immunoreactivity (manifest) (HD grade 1–3)	Not assessed	[87]
<i>Prefrontal cortex</i>			
Volume	↓ Premanifest (VBM)	↓ HdhQ250	[86, 165]
	= Manifest (VBM)		[166]
White matter degeneration	↑ Premanifest (DWI)	Not assessed	[167, 168]
Functional connectivity	↓ Premanifest (fMRI)	↓ BACHD rats	[89, 90, 153]
Activity	↓ Premanifest (fMRI)	↓ R6/2, tgHDrat	[91, 93, 169–171]
Neuronal size	↓ Immunoreactivity (manifest)	Not assessed	[88]
Glial expression	↑ Immunoreactivity (manifest)	Not assessed	[88]
Synaptic integrity	↓ Immunoreactivity (premanifest and manifest)	Not assessed	[172]
Axonal transport	Not assessed	↑ Immunoreactivity (Q175KI)	[173]
Glutamate uptake	↓ Manifest	Not assessed	[174]
BDNF	Not assessed	↓ mRNA levels (R6/2, R6/1)	[142, 175]
Orexin	↓ Immunoreactivity (HD grade 1–4)	Not assessed	[51]
Serotonin	Not assessed	↓ Immunoreactivity (tgHD rat)	[155]

BDNF, brain-derived neurotrophic factor; PSA-NCAM, polysialylated neural cell adhesion molecule; TRH, thyrotropin-releasing hormone; VBM, voxel-based morphometry; fMRI, functional magnetic resonance imaging; DWI, diffusion weighted imaging.

emotional functions are derived from multiple circuit connections involving cortical and subcortical regions of the limbic system, it seems indisputable that limbic system dysfunction can be ascribed to non-motor features of HD. Indeed, volumetric abnormalities in the amygdala have been found to be related to the clinical profile of HD with high level of anxiety and cognitive symptoms occurring before the onset of motor symptoms [77]. This prevailing view of limbic system involvement in HD is also supported by findings from rodent HD models with alterations in volume, reduction in BDNF and serotonin levels reported in several rat and mouse models (see Table 2 for details). The full extent of limbic system pathology in HD is not well understood and more systematic neuropathological postmortem analyses of these structures are warranted.

EFFECTS ON CIRCULATING FACTORS RELATED TO THE HYPOTHALAMUS IN CLINICAL HD

As the hypothalamus is targeted by and regulates circulating (neuro-)endocrine factors in the blood and cerebrospinal fluid (CSF), it became interesting to investigate whether measurements of such circulating factors would be altered in HD as an indication of hypothalamic dysfunction and potentially disease progression. Several studies have been performed using both CSF and blood to measure neuroendocrine factors with often contradicting or negative results [16, 100–104]. The major results are summarized in Table 3. Out of the published results, the increase of cocaine and amphetamine regulated transcript (CART) levels in the CSF is interesting as it may reflect the increased number of CART-immunopositive neurons in the hypothalamus in HD [35, 105]. Also, a subtle change in the hypothalamic-pituitary-adrenal (HPA) axis may be present [106–109]. With the development of more sensitive assays and a better standardization of collected material, it is possible that future studies will reveal other hypothalamic-related factors with relevance for the non-motor features of HD.

LESSONS LEARNT FROM EXPERIMENTAL STUDIES IN HD

There are several animal models expressing different variants of the human mutant huntingtin gene. Several of these transgenic mice display hypothalamic

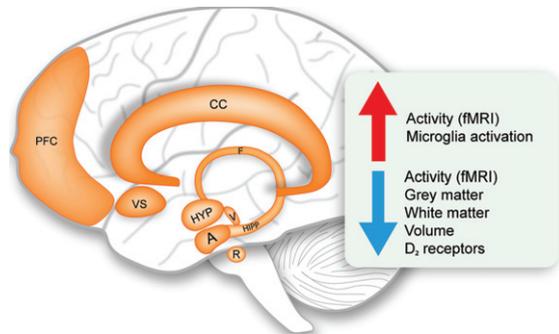


Fig. 2. Limbic system changes in Huntington's disease. Schematic representation of the structures of the limbic system and overview of the main upregulated and downregulated changes (indicated by the arrows). A, amygdala; CC, cingulate cortex; F, fornix; fMRI, functional magnetic resonance imaging; HIPP, hippocampus; HYP, hypothalamus; PFC, prefrontal cortex; R, raphe nucleus; V, ventral tegmental area; VS, ventral striatum.

lamic pathology (Table 4). The R6/2 mouse model shows the highest degree of hypothalamic pathology with reduced number of neurons expressing orexin, oxytocin, vasopressin and VIP [50, 106, 110–113]. Experimental studies provide a complementary approach to the analyses of clinical material as they provide the possibility to study and establish causative relationships. As the hypothalamus receives major input both from the periphery and the rest of the CNS, the question arises as to what extent are the changes detected in this region a direct consequence to the expression of mutant huntingtin in hypothalamic cells. Experiments using recombinant adeno-associated viral (rAAV) vector mediated delivery of different fragments of mutant huntingtin specifically to the hypothalamus have been conducted to address this point. These studies have shown a direct and early effect on the neuronal populations expressing orexin, oxytocin, vasopressin, MCH and BDNF in the hypothalamus, suggesting that the findings made in human postmortem hypothalamic tissue may be directly related to the effects of mutant huntingtin in this area [52, 114, 115]. Other studies have been performed in order to investigate whether the inactivation of mutant huntingtin in the floxed BACHD mouse model using delivery of Cre-recombinase by rAAV vectors or different breeding strategies. rAAV-vector mediated delivery of Cre-recombinase into the hypothalamus has shown a causative link between hypothalamic dysfunction and non-motor phenotypes such as metabolic dysfunction and depressive-like behavior [114, 116]. However, experiments to inactivate mutant huntingtin

Table 3
Summary of main neuropeptide levels in the circulation in both clinical HD and rodent HD models

Neuropeptide	HD stage	HD rodent models	References
<i>Output from the hypothalamus</i>			
ACTH	= Serum, plasma levels (premanifest, HD stage 2-3)	↑ Serum levels (R6/2)	[104, 106, 176]
	↑ Plasma levels (HD stage 1-4)		[126]
AgRP	↑ Plasma levels (premanifest)	Not assessed	[177]
	↓ Plasma levels (manifest)		[177]
CART	↑ CSF levels (early-mid stage HD)	Not assessed	[105]
Cortisol/corticosterone	↑ Plasma, serum levels (HD stage 1-4)	↑ Serum levels (R6/2)	[106, 107, 126, 176, 178-180]
	= Serum levels (premanifest, HD stage 2-3)	= Serum levels (BACHD)	[104, 116]
Estradiol	= Serum levels (premanifest, HD stage 2-3)	= Serum levels (R6/1)	[104, 181]
FSH	= Serum, plasma levels (premanifest, HD stage 2-3)	Not assessed	[104, 176]
GHRF	= Serum levels (premanifest, HD stage 2-3)	Not assessed	[104]
Growth hormone	= Serum and plasma levels (early HD, manifest)	Not assessed	[182, 183]
	↑ Plasma levels (manifest)		[176]
	↓ Plasma levels (manifest)		[177]
IGF-1	= Plasma levels (manifest)	↑ Plasma levels (BACHD, YAC128)	[183, 184]
LH	= Serum, plasma levels (premanifest, HD stage 2-3)	Not assessed	[104, 176]
NPY	↑ CSF levels (early-mid stage HD)	Not assessed	[185]
Orexin	= Ventricular CSF (HD grade 1-4)	= CSF levels (YAC128)	[51, 100, 101, 103, 130]
Prolactin	= Serum, plasma levels (premanifest, HD stage 2-3)	Not assessed	[104, 176, 186, 187]
	↓ Plasma (premanifest and manifest)		[177]
Testosterone	= Serum, plasma levels (premanifest, HD stage 2-3)	↓ Serum levels (R6/1, R6/2)	[104, 139, 176, 181, 188, 189]
	↓ Plasma levels (manifest)	↑ Serum levels (tgHD rat)	[190, 191]
TSH	= Serum, plasma levels (premanifest, HD stage 2-3)	Not assessed	[104, 176, 186]
Vasopressin	= Serum levels (premanifest, HD stage 2-3)	= Serum (BACHD)	[104, 116]
<i>Input to the hypothalamus</i>			
Adiponectin	= Plasma levels (early HD)	↓ Serum levels (R6/2, N171-82Q)	[177, 182, 192]
Amylin	↑ Plasma (premanifest and manifest)	Not assessed	[177]
Ghrelin	= Plasma levels (early HD)	↓ Serum levels (R6/2, N171-82Q)	[110, 182, 192]
	↑ Plasma (manifest)		[177, 183]
GIP	= Plasma (premanifest and manifest)	Not assessed	[177]
Glucagon	↓ Plasma (manifest)	Not assessed	[177]
Glucose	= Plasma levels (manifest)	↑ Serum levels (R6/2, R6/1, N171-82Q)	[106, 183, 192-195]
Insulin	= Plasma levels (HD grade 1-2, manifest)	↑ Serum levels (R6/2)	[106, 177, 183, 193, 194, 196]
		↓ Serum levels (R6/2, N171-82Q)	[192]
Leptin	= Plasma levels (early HD)	↑ BACHD, BAC-225Q, YAC128	[114, 116, 197-199]
	↓ Plasma (manifest)	↓ Serum levels (R6/2, N171-82Q)	[177, 183, 192, 200]
PP	= Plasma (premanifest and manifest)	Not assessed	[177]
PYY	= Plasma (premanifest and manifest)	Not assessed	[177]
Resistin	= Plasma levels (early HD)	Not assessed	[197]

ACTH, adrenocorticotropic hormone; AgRP, agouti-related protein; CART, cocaine and amphetamine regulated transcript; FSH, follicle-stimulating hormone; GHRF, growth hormone-releasing factor; IGF-1, insulin-like growth factor 1; LH, luteinizing hormone; NPY, neuropeptide Y; TSH, thyroid-stimulating hormone; GIP, gastric inhibitory polypeptide; PP, pancreatic polypeptide; PYY, peptide YY.

Table 4
Summary of main hypothalamic findings in HD rodent models

Pathology	HD rodent models	References
Atrophy	↑ R6/2	[110, 131, 201]
Neuronal count	↓ R6/2, N171-82Q	[50, 202]
Upregulation of HPA axis	↑ R6/2	[106]
Circadian rhythm dysfunction	↑ R6/2 (CAG250, CAG 140), BACHD, Q175	[203–207]
Androgen receptor	↓ protein (R6/1)	[139]
Brn2	↓ R6/2	[112]
CART	↓ peptide, immunoreactivity, mRNA (R6/2, HD190QG)	[110, 208]
	↑ mRNA (BACHD)	[116]
Clock genes (mPer2, mBmal1, Cry1, Dbp)	↓ R6/2	[13, 209, 210]
CRH	↓ peptide (R6/2)	[106, 110, 112]
	= mRNA (BACHD)	[116]
GnRH	↓ immunoreactivity (BACHD, R6/2)	[119, 189]
	↓ mRNA (R6/1)	[181]
HAP1 levels	↓ N171-82Q	[202]
MCH	↓ peptide, immunoreactivity, mRNA (R6/2, AAV-hypo)	[52, 110]
	= mRNA (BACHD)	[116]
NPY	↓ mRNA (R6/2, HD190QG, BACHD)	[116, 208]
Orexin	↓ protein, immunoreactivity, mRNA (R6/2, YAC128, AAV-hypo)	[50, 52, 100, 111, 114, 211]
	↑ immunoreactivity (BACHD)	[116]
Orexin 2 receptor	↓ mRNA (BACHD)	[116]
Oxytocin	↓ immunoreactivity, mRNA (R6/2, HD190QG, AAV-hypo)	[112, 114, 208]
	= immunoreactivity, mRNA (BACHD)	[116]
Preprosomatostatin	↓ mRNA (R6/2, HD190QG)	[208]
POMC	↓ peptide, immunoreactivity (R6/2)	[110]
Prodynorphin	↓ mRNA (AAV-hypo)	[52]
Tachykinin receptor 3	↓ mRNA (BACHD)	[116]
TSH releasing hormone	↓ mRNA (R6/2, HD190QG)	[208]
Tyrosine hydroxylase	↓ mRNA (AAV-hypo)	[52]
Vasopressin	↓ mRNA, immunoreactivity (R6/2, HD190QG, AAV-hypo)	[111, 112, 114, 208]
	= immunoreactivity, mRNA (BACHD)	[116]
VIP	↓ Immunoreactivity, mRNA (R6/2)	[113]
VIP receptor	↓ mRNA (R6/2)	[113]

HPA, hypothalamic–pituitary–adrenal; CART, cocaine and amphetamine regulated transcript; CRH, corticotrophin releasing hormone; GnRH, gonadotropin-releasing hormone; HAP1, huntingtin-associated protein 1; MCH, melanin-concentrating hormone; NPY, neuropeptide Y; POMC, proopiomelanocortin; TSH, thyroid stimulating hormone, VIP, vasoactive intestinal peptide.

in specific nuclei or cell populations in the hypothalamus using breeding of the BACHD mice with mice expressing Cre-recombinase under the VMH-specific promoter SF-1, under the PVN-specific promoter Sim-1, or the leptin receptor promoter, did not show any beneficial effects on the psychiatric or metabolic phenotype [117–119]. It would be useful to investigate the involvement of specific hypothalamic circuitries in the development of the early HD phenotype. Further studies using cell specific expression or inactivation of mutant huntingtin in combination with modulating the activity of specific circuitries using chemo- or opto-genetic techniques with analyses of behavioral outputs are likely to give further insight into the causal links between huntingtin-mediated hypothalamic dysfunction and non-motor features of HD.

CLINICAL TRIALS IN HD: FROM A HYPOTHALAMIC PATHOLOGY PERSPECTIVE

According to the clinical trials database (<https://clinicaltrials.gov/>), a total of 124 HD clinical trials have been completed to date worldwide with an additional 46 currently active or in the process of recruiting. These trials have largely focused on restoring motor or cognitive function and very few studies have focused on the non-motor disturbances. Gene therapy approaches to silence the disease-causing mutant HTT protein are currently at the forefront of treatment strategies for HD with the successful completion of the Phase 1/2a IONIS-HTTRx clinical trial (Ionis Pharmaceuticals) using intrathecal administration of antisense oligonucleotides (ASOs) to reduce

both wild-type and mutant HTT [120]. A Phase 3 trial in adult patients with manifest HD (RG6042, Roche Pharmaceuticals) began in April 2019. The hypothalamus is located in a prime position anatomically, surrounding the third ventricle. Although it is likely that intrathecally-delivered ASOs can also target the hypothalamus, preclinical studies using different intrathecally-delivered HTT-lowering agents have not focused on whether there is target engagement in the hypothalamus. With the results of the ongoing HTT-lowering clinical trials, it will be imperative to determine whether intrathecally delivered ASOs reach the hypothalamus and if so, the extent to which this improves the non-motor symptoms and signs. In preparation for future preclinical and clinical trials with HTT-lowering agents, it is worth considering that hypothalamic pathology exists alongside or perhaps even before striatal and cortical pathology. It will be important to further investigate what brain regions need to be targeted to impact upon non-motor features of HD and to pinpoint a treatment timeframe during the life of an affected individual where it is still possible to prevent or reverse symptoms and eventually halt disease progression.

CONCLUSION

The hypothalamus and other areas of the limbic system are part of the brain regions affected in HD. The changes detected mainly in the hypothalamus include effects on factors that regulate motivated behavior, emotions, social cognition, sleep and/or metabolism with reduced levels of orexin, oxytocin, vasopressin, VIP and somatostatin and increased levels of SIRT1 in specific hypothalamic nuclei. Experimental studies in mice have begun to establish causative relationships between expression of mutant huntingtin in the hypothalamus and its effects on neuropeptide expression as well as in the development of depressive-like behavior and metabolic disturbances. Hence, pathology in the hypothalamus may have relevance for the development of non-motor features in HD. Further studies are still needed to investigate the overall extent of these changes in the limbic system and the potential of modulating affected targets for symptomatic effects and/or disease modification in HD.

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CONFLICT OF INTEREST

The authors have no conflict of interest to report.

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