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Huntington's disease and NMDA receptors; a new arena for therapeutic development

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Abstract

Huntington's disease (HD) is an inheritable autosomal dominant neurodegenerative disorder characterized by a clinical triad of progressive choreiform movements (dance-like movements), psychiatric symptoms, and a decline in the cognitive functions. It is caused by a CAG trinucleotide repeat expansion in the HD gene whereas principal pathology of this disease is the loss of striatal and cortical projection neurons. Different experimental animal models have been figured out to assist in understanding the mechanisms involved in the progress of the HD. Extensive researches had been done for testing a large array of neuroprotective agents using animal models which mimic such disorder and to examine the different mechanisms suggested to contribute in the progress of HD. Excitotoxicity in the central nervous system (CNS) results from glutamate exposure for prolonged periods or to excessive concentrations to glutamate. Nmethyl-D-aspartic acid (NMDA) receptor is an ionotropic glutamate receptor responsible for the memory functions and exhibits several features of relevance to neuronal death. Neurotoxicity mediated by NMDA receptor with subsequent loss of striatal neurons contributes in the pathophysiology of HD. The current review focusing on using of NMDA receptor legends as a new strategy in treatment of HD.

Keywords:

Excitotoxicity, Glutamate, Huntington, NMDA, Striatum

1. Introduction

Neurodegenerative diseases of the basal ganglia such as Parkinson's disease (PD) and Huntington's disease (HD) have received considerable concern, particularly in the past few years. In adult-onset HD, patients show a progressive deterioration of motor and cognitive functions that follows three well-defined stages spread over 15-20 years (Harper, 1991). At the initial stage and before the appearance of prominent motor alterations, HD patients are likely to show psychiatric symptoms that include apathy, irritability, depression, and other mood alterations (Reedeker, 2012; Thompson, 2012). Slight motor abnormalities, such as motor tics and jerky voluntary movements, also are likely (Beste et al., 2009). The second stage is characterized by a tragic increase in involuntary movements, which become generalized, abrupt, and uncontrolled. Deterioration of daily activities such as walking, eating, speaking, and swallowing occurs as choreic movements become more prominent. In some cases, bradykinesia may exist concomitantly with the choreic movements that become apparent in the third stage (Thompson et al., 1988). Cognitive abilities progressively decline and finally end with dementia. Another hallmark of the second stage is the loss of body weight (Marder et al., 2009). Overall health progressively worsens and by the third stage, nearly 10-15 years after diagnosis, the choreic movements are replaced by bradykinesia and rigidity. Death becomes abutting, and the most common causes are pneumonia and heart disease.

2. Genetic basis & Neuropathology

Huntingtin (HTT) was the first diseaseassociated gene to be molecularly mapped to a human chromosome (Gusella et al., 1983). Ten years later, the DNA sequence has been identified and the precise nature of the HDassociated mutation in HTT was determined (MacDonald al., 1993). et The HTT gene contains a region where the triplet nucleotide (CAG) is repeated several times. These CAG repeats generate a series of glutamine residues known as a polyglutamine tract (polyQ) and the number of CAG repeats present in the HTT gene determines whether an individual will have HD or not. Individuals less than 35 CAG repeats will be unaffected; if repeats increased up to 36-39 repeats, the individual will be at increased risk for HD; and individuals with 40 or more

repeats will definitely manifest disease phenotypes (Bates, 2005).

A rough inverse correlation between the number of CAGs and the age of HD onset occur (Langbehn et al., 2010). CAG repeats ranging from 50 to 200 (the highest number reported) may precipitate HD onset in childhood or teenage years that is characterized by psychiatric disturbances, accelerated mental and physical deterioration that culminates in death as soon as 5–10 years after onset (Quarrell et al., 2012).

Although the exact cause of neuronal death in HD remains unknown, it has been postulated that proteolysis of mutant huntingin plays a role in disease pathogenesis resulting in an abnormal and toxic amino acid (N) terminal fragment that forms protein aggregates in neurons. Intra-cellular aggregates of mutant HTT are found in both the cytoplasm and nucleus (Vonsattel et al., 1985). In neurons, mutant HTT aggregates are also located in axon terminals, where the presence of aggregates has been correlated with decreased density of synaptic vesicles and decreased release of glutamate in striatum of HD mice (Gafni et al., 2004; Finkbeiner and Mitra, 2008) These aggregates sequester a variety of important cellular proteins leading to disruption of protein can be readily detected homeostasis and throughout the course of the disease. Some evidence has outlined that direct interactions between mutant huntingtin and transcription factors may be the proximal defect that leads to transcriptional dysfunction (Tunez et al., 2010). This then results in a cascade of compensatory and damaging events as mitochondrial dysfunction, energy depletion, oxidative stress, DNA damage and others that might play a role in neuronal death (Pringsheim et al., 2012).

The most prominent cell loss in HD occurs in cerebral cortex and striatum, the forebrain regions in which cortical pyramidal neurons (CPNs) and striatal medium spiny neurons (MSNs) are the most affected (Tunez et al., 2010).

3. Animal models of HD

3.1. Assessing the validity of animal models of human disease

The applicability of a given animal model of human disease is often judged based on three broad measures of validity: the animal model's *construct validity* (refers to how closely the animal model mimics the underlying pathogenic lesion that provokes the disease in humans), *face validity* (relates to the extent to which the animal model develops the symptoms and phenotypes associated with the human disease) and *predictive validity* (indicates how closely improvements in response to treatment in the animal model parallel, or predict, improvements in patients). For diseases that lacks an effective treatment, such as HD, assessment of the predictive validity of an animal model is currently impossible (Pouladi et al., 2013).

3.2. Models for HD:

A number of animal models of HD have been available for more than 30 years (Brouillet et al., 1999) in an effort to understand how the mutation causes the symptoms and pathology observed in patients in order to observe the process of cell death in a simpler, more controlled circumstances than the human disorder.

3.2.1. Toxin Models of HD:

3.2.1.1. Neurotoxin-mediated lesions in animals:

Early experimental models of HD relied on direct introduction of excitatory agonists into the central nervous system (Rothman, 1984). Kainic acid is an excitatory glutamate-type neurotoxin, which produces degeneration of striatal GABAgenic projection neurons with preservation of striatal afferents (McGeer and McGeer, 1976). Kianic acid was also found to irreversibly reduce glutamic acid decarboxylase (GAD) but do not produce the entire constellation of histological changes observed in HD (Isacson et al., 1985; Hantraye et al., 1990).

Unlike kainic acid, quinolinic acid shows differential sparing of striatal neurons (Lehrmann et al., 2008). Quinolinic acid damages both GABAergic and substance P-containing neurons, with relative sparing of cholinergic neurons, that latter of which are known to be spared in HD (Ferrante et al., 1985; Ferrante et al., 1987).

The behavioural deficits in the excitotoxin rodent model of HD include locomotor hyperactivity and poor performance in mnemonic and cognitive tests (Koutouzis et al., 1994) but doesn't include dyskinesias or chorea like movements. Thus, the rodent excitotoxin model can't be used to study the progression of the disease or to replicate the widespread neuropathology observed in the human.

3.2.1.2. Defective energy metabolism toxin models:

Impaired cellular energy maybe involved in the degenerative process in HD (Albin and Greenamyre, 1992) due to reduced ATP stores that results in membrane depolarization, removal of magnesium from the NMDA linked calcium channel and subsequent excitotoxic injury (Novelli et al., 1988).

3-Nitropropionic acid (3-NP) has been associated with neurological illness in animals and humans (Ludolph et al., 1991). 3-NP is an irreversible inhibitor of succinate dehydrogenase (SDH) that inhibits both the Krebs cycle and complex II activity of the electron transport chain. Moreover, striatal atrophy is apparent with systematic administration of low doses of 3-NP (Beal et al., 1993; Brouillet et al., 1993).

3.2.2. Genetic models of HD:

In particular, genetic animal models of HD have been an accurate tool of elucidation of the progression of behavioral and physiological alterations, which had not been possible using classic neurotoxin models. Genetic models also provide an opportunity to test promising treatments and explore their potential ability to cure the disease (Ferrante et al., 2009).

3.2.2.1. R6/2 transgenic mice:

line of transgenic mice The R6 (Mangiarini et al., 1996) remains one of the most widely used models because it offers many advantages. Particularly, R6/2 mice (expressing exon 1 with about 150 CAG repeats) manifest a very aggressive, rapidly progressing phenotype, similar to the juvenile form of HD in humans. They display overt behavioural symptoms as early as 5-6 weeks of age and die at about 15 weeks. Pathophysiological alterations include the formation of nuclear inclusions (Davies et al., 1997), which were later also shown to be present in human HD brains and can be observed in the presymptomatic stage, particularly in the striatum and CA1 region of the hippocampus (Morton et 2000). There are also changes al., in neurotransmitter receptor expression (Cha et al., 1998; Ariano et al., 2002) and altered signaling mechanisms (Bibb et al., 2000; Luthi-Carter et al., 2000; Menalled et al., 2000). Many of these alterations are correlated with motor (Carter et al., 1999) and learning impairments on a number of cognitive tasks (Lione et al., 1999), as well as deficits in synaptic plasticity in the hippocampus before an overt phenotype appears (Murphy et al., 2000) and in the striatum of symptomatic animals (Kung et al., 2007).

3.2.2.2. R6/1 transgenic mice:

The R6/1 transgenic mice were developed at the same time as the R6/2 transgenic mice and express exon 1 of the human HD gene with approximately 116 CAG repeats. The R6/1 transgenic mice have not been as well studied as the R6/2 transgenic mice (Mangiarini et al., 1996). The R6/1 line has a later age of onset and a slower disease progression. They can live beyond 12 months (Naver et al., 2003).

3.2.2.3. N171-82Q transgenic HD mice:

N171-82Q mice have an N-terminal fragment of huntingtin incorporating both exon 1 and exon 2 of the huntingtin gene, with 82 polygulatamines (Schilling et al., 1999). The commercially available line has a life span range from 130-180 days (Yu et al., 2003).

3.2.2.4. Murine huntingtin homologue knock-in mice:

Knock-in models also contributed to understanding of HD. The major advantage of these models is that they express full-length mutant htt in its native genomic context. Several models have been generated and the difference is mainly in the number of CAG repeats (from 48 to 200) (Heng et al., 2007; Heng et al., 2009) (White et al., 1997; Levine et al., 1999; Wheeler et al., 2000; Lin et al., 2001). In the knock-in models early overt behavioural changes are often subtle, but sensitive and careful testing demonstrate abnormalities as early as 1–2 months of age (Menalled et al., 2002; Menalled et al., 2003).

A more severe phenotype develops as the mice age beyond one year (Hickey et al., 2008). A consistent feature of knock-in mice is the presence of nuclear staining and microaggregates very early in the course of the disease. By contrast, nuclear inclusions are observed only in older mice (10–18 months) (Menalled and Chesselet, 2002), and loss of MSNs occurs at about 2 years (Hickey et al., 2008).

A number of lines have been made:

Hdh/Q 72-80: with 72 to 80 CAG repeats (Alexi et al., 1998).

HdhQ111: with 109 to 111 CAG repeats (Brouillet et al., 1998).

CAG140: with 140 CAG repeats (Naver et al., 2003).

CAG150: with 150 CAG repeats (Bizat et al., 2003).

3.2.2.5. Full length human HD gene transgenic mouse models:

A yeast artificial chromosome mouse model of HD with the entire human huntingtin gene containing 126 CAG repeats as well as flanking segments that might involve regulatory elements. Motor abnormalities develop consisting of initial hyperactivity followed by difficulty walking and then hypokinesia (Hodgson et al., 1999; Slow et al., 2003).

4. Mitochondria in HD

Mitochondrial function is impaired in HD and this occurs early in the disease process. The striatum might be particularly vulnerable to mitochondrial defects through multiple mechanisms involving molecular factors, as, dopamine and D_2 receptors and/or glutamate and NMDA receptors, which are selectively present in this brain in high concentrations and eventually play a role in the selective degeneration of striatal neurons expressing mutant Htt (Damiano et al., 2010).

Intrastriatal injection of malonate (a reversible inhibitor of succinate dehydrogenase) supported the hypothesis that impairment of mitochondrial function plays an important role in the pathogenesis of HD (Beal et al., 1994). Also, mitochondrial dysfunction is evident in two well-established HD mice models; the 150/150Q mutant huntingtin knock-in mice (Lin et al., 2001), and the R6/2 mice (Mangiarini et al., 1996).

The formation of reactive oxygen species (ROS) and the imbalance between oxidant and antioxidant systems due to mitochondrial dysfunction usually associated with cellular damage and neuronal death that plays a crucial role in the neurodegenerative process of HD (Beal, 1995; Banoei et al., 2007; Browne, 2008; Gil and Rego, 2008; Lim et al., 2008, Kalonia, 2009; Kumar et al., 2009; Tasset et al., 2009; Tunez and Santamaria, 2009; Kumar and Kumar, 2010).

5. Glutamate induced excitotoxicity

Glutamate is the main excitatory neurotransmitter in the vertebrate central nervous system, it's essential for a wide variety of physiological processes, such as integrative brain function and neuronal cell development and it's also implicated in neuronal cell death and has been postulated to play an important role in the pathogenesis of different degenerative disorders (Meldrum, 2000; Koller and Cersosimo, 2004; Rego and de Almeida, 2005; Yi and Hazell, 2006; Corona et al., 2007; Hazell, 2007; Mamelak, 2007; Farooqui, 2008).

The uncontrolled release of glutamate is known to induce a pathological process named excitotoxicity (Olney, 1994). Excitotoxicity is one of the most extensively studied processes of neuronal cell death, and it plays an important role in many CNS diseases, including ischemia, trauma, and neurodegenerative disorders (Dong et al., 2009).

Many electrophysiological data support the view that abnormal glutamatergic transmission in the corticostriatal pathway is a prominent aspect of striatal degeneration in HD (Cepeda et al., 2007). Also, decortication which removes the central striatal glutamate, protects against striatal degeneration in R6/2 mice model of HD (Stack et al., 2007). As shown in (**Figure 1**), there are at least three different types of ionotropic glutamate receptor, N-methyl-D-aspartate (NMDA) receptor, alpha-amino-3-hydroxy-5-methyl-4-

isoxazolepropionic acid (AMPA), and kainate receptors. and eight distinct metabotropic glutamate receptors (mGluRs), which are divided into three subgroups based on sequence homology and G protein coupling specificity (Pin and Duvoisin 1995, Conn and Pin 1997, Dhami and Ferguson 2006). Group II and Group III of mGluRs are mainly localized presynaptically (autoreceptors) and have become a target of potential therapeutics for various neurological diseases and conditions, because of the ability of agonists to suppress glutamate release (Kew et al., 2001; Schoepp, 2001).

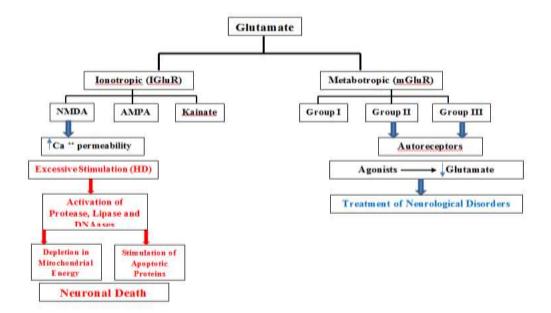


Fig.1. Types of glutamate receptors and their involvement in excitotoxicity and treatment of neurological disorders.

6. Role of NMDA receptors in HD

NMDA receptors are tetrameric complexes of two subunit 1 (GluN1) (NR1) with two GluN2 (NR2) and/or GluN3 (NR3) subunits. The differences in receptor channel properties, pharmacology, and temporal and spatial brain distribution are determined by these subunits (Cull-Candy and Leszkiewicz, 2004). Considerable literature suggests that activation of extrasynaptic NMDA receptors promotes neuronal death, while activation of intrasynaptic NMDA receptors promoting neuronal survival (Hardingham et al., 2002, Leveille et al., 2008). Other literature suggests that NR2B containing NMDA receptors are excitotoxic mediators, while NR2A containing NMDA receptors promote neuronal survival (Hansson et al., 2001). Striatal neurons express both NR2A and NR2B containing NMDA receptors and it is likely that medium spiny extrasynaptic NMDA receptors preferentially contain NR2B subunits (Stack and Ferrante, 2007).

NMDA receptors activation requires presynaptic glutamate release coupled with postsynaptic depolarization to allow ion flux through the channel. Also, overstimulation of NMDA receptors results in activation of proteases, DNAases, mitochondrial failure, and/or signalling to a variety of apoptotic proteins, leading to neuronal death (Lipton and Rosenberg, 1994; Arundine and Tymianski, 2003). Thus, NMDA receptors are ideal candidates for contributing to early neuronal degeneration in HD.

Several lines of evidence suggest that abnormal activation of NMDA receptors and subsequent excitotoxicity aggravates striatal neuronal loss and plays important role in HD pathogenesis (Wojda et al., 2008). Previous studies have shown that intrastriatal injection of NMDA receptor agonists, including NMDA and quinolinic acid laboratory animals produces in neurochemical, neuropathological, and behavioural changes characteristic to HD (Laforet et al., 2001). Also, increased NMDA receptor mediated currents and enhanced susceptibility to exogenous NMDA agonists are described in murine genetic models of HD (Fan and Raymond, 2007).

Also, NMDA receptor-mediated excitotoxicity and cell death were enhanced in many non-neuronal cell lines expressing mutant Htt and in striatal neurons from transgenic mice models of HD, for example; data obtained from experiments using YAC72 mouse model (Fan and Raymond, 2007) showed overexpression of the synaptic NMDA receptor subunits of the striatal neurons, which would aid to understand the increased current and toxicity, and another example where mutant Htt increases the sensitivity of neurons to excitotoxicity induced by stimulation of NMDA receptors bearing NR2B subunits (Zeron et al., 2001; Zeron et al., 2002). The specificity of these effects for the NR2B-subtype of NMDA receptors may explain partly the selective degeneration of certain neurons in HD.

The extent of sensitivity to excitotoxic insult is not correlated to NMDA receptor current. The R6/2 transgenic mouse model of HD exhibits an increased NMDA receptor-mediated current and a decreased sensitivity to quinolinic acid

excitotoxic insults (Zeron et al., 2002; Zeron et al., 2004; Li et al., 2003; Li et al., 2003).

Also, studies showing that MSNs are more vulnerable to NMDA receptors mediated toxicity compared to other striatal neuronal types led to the hypothesis that enhanced NMDA receptors activation could contribute to selective striatal degeneration in HD. For example YAC72 and YAC128 mice showed increased striatal lesion size after quinolinic acid injection compared to controls (Graham et al., 2009). Remarkably, R6/1 and R6/2 mice exhibit wide spread brain atrophy, without selective striatal neuronal degeneration, whereas YAC transgenic HD mice show mild but significant striatal neuronal loss with lesser losses of cortical neurons and no significant atrophy in other brain regions (Slow et al., 2003). These data suggest that protein context is important for accurately modelling selective neuronal degeneration, and that the YAC and other fulllength mutant Htt models may be better suited for study of the earliest molecular mechanisms underlying striatal neuronal susceptibility.

To be mentioned, striatal NMDA receptormediated responses are enhanced in HD transgenic mice prior to onset of motor deficits and neuronal loss (Li et al., 2004; Andre et al., 2006; Milnerwood and Raymond, 2007), and this consistent with studies showed that intrastriatal injection of quinolinic acid produces a significantly larger lesion in YAC128 and R6/1 models prior to symptoms onset (Graham et al., 2009).

7. Effect of NMDA receptor antagonists on treatment of HD

NMDA receptor antagonists describe a category of compounds that functionally inhibit or deactivate NMDA receptor activity. They can act broadly or specifically at various sites on the NMDA receptor including the agonist binding domains, allosteric sites, and the ion channel pore. This category of compounds has a potential use in any disease that results from glutamate-induced excitotoxicity ranging from epilepsy to neurodegenerative disorders like HD (Leveille et al., 2008).

Depending on the excitotoxic nature of NR2B containing NMDA receptors especially those present in striatal medium-sized spiny neurons, blockage of this type of receptors with selective antagonist (subunit specific compounds) is a rational approach to neuroprotection in HD. NR2B-selective antagonist (remacemide) ameliorates the phenotype of the R6/2 transgenic model of HD with improved therapeutic efficacy and minimum side effects (Stack et al., 2007). Also, clinically available antagonists such as memantine or amantadine may be candidates for large clinical trials of HD, but some trials demonstrated little therapeutic value due to the adverse effects of synaptic NMDA receptor blockade (Fan and Raymond, 2007).

balance between synaptic and The extrasynaptic NMDA receptor activation appears to determine whether receptor activation is beneficial or detrimental (Hardingham et al., 2002). It has been shown that treatment of YAC128 mice with low dose of memantine, which preferentially blocks extrasynaptic NMDA receptors, was ameliorative and therapeutically effective. While at higher doses, memantine treatment exacerbated striatal degeneration in YAC128 mice, most probably, due to loss of selectivity and the blockade of both synaptic and extrasynaptic NMDA receptors, obviating the benefits of inhibiting extrasynaptic NMDA receptors and possibly exacerbating mutant huntingtin toxicity by inhibiting the neuroprotective effects of synaptic NMDA receptor activation (Okamoto et al., 2009).

Another study (Tallaksen-Greene et al., 2010) demonstrated lack of efficacy of selective NR2B-antagonists such as; ifenprodil, RO25,6981 and CP101,606 in the R6/2 model of HD due to use of them in high doses in an effort to obtain maximal effects. It is plausible that this approach resulted in blockade of pro-survival signaling effects of synaptic NMDA receptors, counteracting any beneficial effects of reducing the excitotoxic effects of extrasynaptic NMDA receptors.

Therefore, targeting extrasynaptic NMDA receptors specifically is a very difficult issue and determination of appropriate dose of any selective NR2B antagonist for neuroprotective trials in HD subjects will be crucial. Dose selection may require some type of human pharmacodynamic biomarker of neuroprotective NMDA receptor blockade which consider a formidable obstacle.

8. Conclusion

HD is an autosomal dominant neurological disorder. Clinical course of the disease involves motor, cognitive and behavioural symptoms. Several animal models have been used to describe the underlying mechanisms of disease progression. One of these mechanisms is the NMDA receptors mediated excitotoxicity. NMDA receptor is an ionotropic glutamate receptor that regulates Ca^{+2} efflux and contributes to the progress of HD. Thus, the use of NMDA receptor antagonists especially those which are selective to extrasynaptic NR2B containing NMDA receptors may be of therapeutic interest to treat HD, but selection of appropriate dose is a must.

Conflict of Interest

The authors report no declaration of conflict of interest.

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