Rectifying rectifier channels in Huntington disease

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Commentary to: Tong X, Ao Y, Faas GC, Nwaobi SE, Xu J, Haustein MD, Anderson MA, Mody I, Olsen ML, Sofroniew MV, et al. Astrocyte Kir4.1 ion channel deficits contribute to neuronal dysfunction in Huntington's disease model mice. Nat Neurosci 2014; 17:694-703; PMID:24686787; http://dx.doi.org/10.1038/nn.3691 Neuronal abnormalities in neurodegenerative disorders such as Huntington disease, Alzheimer disease or Parkinson disease have been the primary focus of decades of research. However, increasing evidences indicate that glial cells and more specifically astrocytes could be as important players as their big brother. It is now particularly evident in Huntington disease where astrocytal potassium channels have emerged as a likely key factor in the pathogenesis of the disease.

Huntington disease (HD) is an inherited neurodegenerative disorder, caused by an autosomal dominant mutation in the Huntingtin gene (HTT).¹ In 1872, Gorge Huntington first described the medical history of several generations of a family experiencing similar symptoms, suggesting that these symptoms must be linked.² The disease was later named after him and is characterized by alterations in personality, cognition, and motor control, associated with neuronal dysfunction and atrophy of the striatum, the substantia nigra, the CA1 region of the hippocampus, and in a lesser extent other brain regions.³ HD is much more common in Europe, North America, and Australia than it is in Asia, with a prevalence of 5-7 and 0.4 affected individuals per 100.000, respectively. This difference is mainly explained by the existence of various huntingtin gene haplotypes.^{3,4} The cause of HD is a wellknown expansion of CAG trinucleotides located at the N-terminal region of the huntingtin protein (HTT).5 Mutated huntingtin (mHTT) tends to aggregate and form intracellular accumulations, a

common cellular phenotype also commonly observed with other neurodegenerative diseases.^{6,7} The physiological role of HTT as well as the pathological mechanism of the disease is largely unknown. However, there are a few known interaction partners that link HTT to vesicular transport, cytoskeletal organization, presynaptic signaling, and anti-apoptotic factors.8 The initial observation that mHTT accumulates in striatal astrocytes^{9,10} and the fact that ion channels have been implicated in neurodegenerative diseases earlier¹¹ lead Tong et al.¹² to investigate the functional implication of glial cells in the pathophysiology of HD. Using mouse models of HD, the authors demonstrate that striatal astrocytes containing mHTT present altered Kir4.1 channel activity and aberrant extracellular potassium homeostasis rendering striatal neurons hyperexcitable, which possibly account for HD motor symptoms.

Intracellular inclusions of mHTT were found in striatal astrocytes of R6/2 mice at a symptomatic stage of HD, before any signs of astrogliosis. Astrogliosis is defined as an increase in astrocyte population due to destruction of nearby neurons, also often observed in brain trauma, stroke, ischemia, and neurodegenerative diseases. Indeed, while old R6/2 mice (P104-P110) with established neurodegeneration and striatal atrophy present an increase in Glial Fribrillary Acidic Protein signal (GFAP, an astrocyte marker), GFAP reactivity remained unaltered in symptomatic R6/2 mice (P60-80) indicating that HD symptoms start before any astrogliosis.



Figure 1. A putative model linking astrocytal Kir4.1 channel and glutamate homeostasis in Huntington disease. In normal condition, glutamate released in the synaptic cleft is rapidly removed by astroctal EAAT transporters that serve o terminate the excitatory signal and preventing neuronal excitotoxicity. Under pathological condition such as Huntington disease, decreased Kir4.1 channel activity leads to aberrant K⁺ homeostasis and transmembrane K⁺ gradient disturbing EAAT activity and astrocytal glutamate uptake. Accumulation of glutamate in the synaptic cleft causes in turn neuronal hyperexcitability and in the long-term cellular toxicity, possibly by activation of extrasynaptic NMDA receptors.

In addition, pSTAT3 (Signal Transducer and Activator of Transcription 3), a transcription regulator usually increased during astrogliosis was found stable in symptomatic R6/2 mice compare with wild type animals.

Using another mouse model of HD, the authors found that striatal astrocytes from HDR6/2 mice show altered electrical properties including a decreased membrane conductance and depolarized resting membrane potential. In addition, the inward-rectifying Kir4.1 potassium (K⁺) channel activity was found significantly reduced. Kir4.1 channel is one of the predominant astrocytal K⁺ channels13-15 that contribute to establish the resting membrane potential of the cells by passing more easily K⁺ ions inside the cell than outside, and also responsible for buffering extracellular K⁺ in the brain. Mutations in Kir4.1 channel are associated with some forms of epilepsy and ataxia.¹⁶ Patch-clamp recordings on astrocytes isolated from R6/2 and Q147 HD mouse indicate that the Ba2+ sensitive K⁺ current, i.e., the K⁺ current conducted by Kir channels, is significantly

reduced at the symptomatic stage but not at a pre-symptomatic time. Reduced Kir currents are most likely the result of either an alteration in Kir channels expression at the plasma membrane or from a direct alteration of channel activity since no change in Kir4.1 mRNA was noticed in R6/2 striatal astrocytes. Consistent with an altered Kir4.1 activity, extracellular potassium concentration was found increased from 1.5 mM to 3 mM, and similar increase in K⁺ concentration applied on striatal slice from wild type animals was sufficient to make neurons hyperexcitable. Finally, viral delivery of Kir4.1 channel into R6/2 mouse striatum was found sufficient to rescue astrocyte electrical properties and normalize extracellular K⁺ level, while only some of the motor symptoms were improved in mice, indicating that Kir4.1 channels contribute to HD symptoms but most likely are not sufficient to fully explain the pathogenesis of the disease. Hence, a decreased expression of the glutamate transporter EAAT (Excitatory Amino-Acid Transporter), which play a critical role in removing glutamate from

the synaptic cleft and thus terminating synaptic transmission¹⁷ was also observed in R6/2 striatum.

While implication of Kir4.1 channel is unambiguous, the cellular mechanisms by which alteration in channel activity contributes to HD symptoms remain intriguing. It is possible that Kir4.1-dependent alteration of neuronal excitability could involve glutamate signaling as it has already been proposed.9,10 Indeed, while activity of the EAAT relies on extracellular Na⁺, it also strongly depends on transmembrane K⁺/ H⁺ concentration gradient.¹⁸ Given the decreased activity of Kir4.1 channel and subsequent increase in extracellular K⁺ concentration in HD mouse models, it is reasonable to think that activity of the glutamate transporter could be altered in those animals. Decrease in EAAT expression, combined with a decreased activity of the transporter could in turn let the extracellular glutamate building up and cause neuronal hyperexcitability and in the long-term cellular toxicity, possibly by acting on extrasynaptic NMDA receptors that have been particularly implicated in

glutamate-associated toxicity¹⁹ (see also Fig. 1). Further investigations to assess extracellular glutamate homeostasis in those HD animal models will certainly provide important insights into the cellular and molecular mechanisms by which astrocytal Kir4.1 channels contribute to HD symptoms. Moreover, considering that an alteration in Kir4.1 channel expression and altered K⁺ homeostasis was also reported in animal mouse models of Alzheimer disease,²⁰ Kir4.1 channel appears as a common underlying mechanism of those neurological disorders and possibly opens interesting new therapeutic avenues.

Disclosure of Potential Conflicts of Interest

No potential conflicts of interest were disclosed.

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