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Cannabinoids: Do they have the potential to treat the symptoms of multiple sclerosis?

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Abstract

This article reviews the role of cannabinoids in inhibiting neurodegeneration in models of multiple sclerosis (MS). MS is a chronic, debilitating disease of the central nervous system (CNS), induced by autoimmunity-driven inflammation that leads to demyelination and thus disconnection of the normal transmission of nerve impulses. Despite the use of an array of immune modulating drugs that restore blood brain barrier function, disability continues in patients concomitant with the loss of axons in the spinal cord. MS patients therefore suffer neuropathic pain, spasticity and tremor. Anecdotal evidence suggests that MS patients using cannabis, though illegal, achieve symptomatic relief from neuropathic pain and spasticity associated with MS. The discovery of the endogenous cannabinoid (endocannabinoid) system that naturally exists in the body and which responds to cannabinoids to exert their effects has aided research into the therapeutic utility of cannabinoids. The endocannabinoid system consists of two G-protein coupled receptors cannabinoid receptor type-1 (CB₁) and CB₂.

CB₁ is mainly expressed in the CNS and CB₂ is predominantly found in leukocytes, while an increasing number of potential ligands and endocannabinoid degradation molecules are being isolated. Several studies have highlighted the involvement of this system in regulating neurotransmission and its ability to prevent excessive neurotransmitter release, consistent with a capacity to provide symptomatic relief. In summary, antagonism of the CB₁ receptor pathway contributes to neuronal damage in chronic relapsing experimental allergic encephalomyelitis (EAE) and suppresses tremor and spasticity. The addition of exogenous CB₁ agonists derived from cannabis also afforded significant neuroprotection from the consequences of inflammatory CNS disease in EAE and experimental allergic uveitis models. Although clear neuroprotective benefits of cannabinoids have been demonstrated, the unwanted psychotropic effects need to be addressed. However, manipulating the endogenous cannabinoid system may be one way of eliciting beneficial effects without some or all of the unwanted side effects.

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Key words: Multiple sclerosis; Axonal damage; Neurodegeneration; Neuroprotection

Core tip: Multiple sclerosis (MS) is an inflammatory demyelinating disease of the central nervous system and causes disability, neuropathic pain, spasticity and tremor in affected patients. Although illegal, users of cannabis report relief from pain and spasticity, probably due to the endogenous cannabinoid system that exists. Cannabinoid receptor type-1 (CB₁)-deficient mice accrue greater levels of neurodegeneration and poorly tolerate inflammatory and excitotoxic insults after immune attack in a model of MS, experimental allergic encephalomyelitis. Treatment of animals affected by experimental allergic uveitis (EAU) with CB₁ agonists also provided significant neuroprotection from the con-

sequences of EAU, suggesting that cannabinoids may slow down neurodegeneration in MS.

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BIOGRAPHY

Dr Zubair Ahmed received his PhD from University College London, London, United Kingdom. He completed his first postdoctoral training period at the Institute of Neurology in London before moving to the University of Birmingham to continue his second postdoctoral training period. In 2007, he was awarded an Research Councils United Kingdom Academic Fellowship, the remit of which was to develop an independently funded research group before being promoted to Lecturer in 2011. He is currently a Senior Lecturer in Neuroscience. His research interests covers numerous aspects of the molecular biology of central nervous system (CNS) axon regeneration and degeneration and his most notable contribution is to the understanding of the role of caspase-2 in apoptosis of retinal and dorsal root ganglion neurons, the contribution of an endogenous mechanism for receptor shedding, the identification of growth factors capable of promoting retinal ganglion cell survival and axon regeneration, identification of alternative neuronal receptors capable of interacting with key molecules in blocking CNS axon regeneration and the observations that cannabinoids inhibit neurodegeneration in models of multiple sclerosis (MS). His curriculum vitae lists over 50 peer-reviewed publications, 2 book chapters and numerous presentations at national and international meetings.

INTRODUCTION

MS is an inflammatory demyelinating disease of the CNS and results in disruption to the normal transmission of nerve impulses due to lesions in the CNS^[1,2]. Despite immune modulating drugs that reduce blood brain barrier dysfunction, disability often continues in patients and suggests that neurodegenerative changes are key to the progression of disease^[3-5]. MS patients thus display spasticity, neuropathic pain associated with neuroinflammation, excitotoxicity and chronic neurodegeneration. It is also established that axonal/neuronal loss, which occurs early in the disease process, is an important contributor of permanent disability and is often associated with active inflammation. Disability only becomes evident when normal compensatory mechanisms are exhausted, while demyelinated axons are left vulnerable to further damage by electrical activity^[6,7]. Therefore identifying neuroprotective strategies are a key goal in the fight against MS.

Although the *Cannabis sativa* plant has been used for centuries as a medicinal preparation to relieve the

symptoms of inflammatory and neuropathic disorders, its use is illegal^[8]. However, anecdotal accounts from MS patients indicated that cannabis might offer symptomatic relief of pain and spasticity associated with MS^[9]. Plant-derived “cannabinoids” have provided important insights into the biology of cannabis and have led to a multitude of clinical trials using cannabinoids to control pain and spasticity in MS patients^[10]. The main active ingredient of *Cannabis sativa* was defined in 1964 as (-)-trans-delta-9-tetrahydrocannabinol [Δ^9 -THC or dronabinol (international non-proprietary name)]^[11]. Δ^9 -THC is not only responsible for the majority of the pharmacological actions of cannabis but also its psychoactive effects. Numerous other cannabinoids and phytochemicals also exist in the cannabis plant including cannabidiol (CBD) which is non-psychoactive and is not a cannabinoid receptor agonist^[12].

CLINICAL TRIALS OF CANNABIS IN MS

Although many patients self-medicated with cannabis, there were few clinical studies that demonstrated reliable clinical evidence of the benefits of cannabis use in MS. Some of the first few studies on the effect of Δ^9 -THC were not encouraging since small sample sizes were used and Δ^9 -THC had no effect on objective measures, despite patients reporting subjective improvements in the symptoms of MS^[13]. For example, the first systematic, placebo-controlled trial with Δ^9 -THC and a *Cannabis sativa* plant extract, given to 16 MS patients with severe spasticity, showed no effect of the cannabinoids on spasticity^[14]. A much larger trial involving 660 MS patients receiving Δ^9 -THC or natural Cannabis oil or a placebo, the mean reported effect on spasticity was not significantly different between control and treatment groups^[15]. However, patient-reported spasticity was reduced, agreeing with earlier studies that showed subjective improvements in spasticity related to MS^[14,15]. However, in a 12-mo follow up study of 657 patients, Δ^9 -THC was reported to have a significant effect on the objective measures of spasticity^[16]. Several later studies have also shown variable results depending on cannabis preparation, dosing regime and patient numbers, throwing into doubt the use of clinical measures of spasticity.

At present, the large number of clinical trials in MS have not clarified whether cannabis is beneficial in MS but have thrown up questions about the rating of “spasticity”, route of delivery, source and dosing regime^[10]. Despite these reservations, self-medicated use of cannabis continues. In a postal questionnaire involving the responses received from 110 MS patients in the South of England, 43% confirmed their use of cannabis with 68% of these patients specifically using cannabis to relieve the symptoms of MS^[17]. Patients affirmed that their main reason for choosing to self-medicate with cannabis was due to pain and spasms, while a small proportion used cannabis for sleep related problems^[17]. In a further study, over 90% of a cohort of 112 MS patients based in the United Kingdom and United States declared that

self-medication with cannabis improved nocturnal pain, spasms and muscular pain^[18].

A recent study showed that of the 572 MS patients enrolled in a clinical study, 272 (47.6%) responded to Sativex, an oral-mucosal spray that contains Δ^9 -THC and CBD, treatment within 4 wk with a response that was defined as > 20% decrease in spasticity^[19]. A second phase of this study demonstrated that the cannabis extracts significantly reduced spasticity and the frequency of spasms while improving sleep quality over an extended 12-wk period, compared to placebo controls^[19]. This has led to approval of the use of Sativex in the treatment of MS-related spasticity. These studies demonstrate the potential beneficial roles of cannabis use in MS patients, although in general its supply and use remains illegal.

ENDOCANNABINOID SYSTEM

The discovery of the endogenous cannabinoid (endocannabinoid) system that naturally exists in the body and responds to cannabinoids to exert their effects has aided research into the therapeutic utility of cannabinoids. This has fuelled the search for alternative modes of delivery into the human body, rather than the traditional method of “smoking”. The endocannabinoid system consists of at least two families of lipid signalling molecules (the N-acyl ethanolamines and the monoacyl-glycerols), multiple enzymes in the biosynthesis and degradation of these lipids, as well as two G-protein coupled receptors [cannabinoid receptor type-1 (CB₁) and CB₂] (reviewed by^[10]). CB₁ is mainly expressed in the CNS while CB₂ is predominantly found in leukocytes, while an increasing number of potential ligands and endocannabinoid degradation molecules are being isolated^[20]. Several studies have highlighted the involvement of this system in regulating neurotransmission and its ability to prevent excessive neurotransmitter release, consistent with a capacity to provide symptomatic relief in MS^[21,22].

Endocannabinoids are produced on demand and are retrogradely transported across the postsynaptic membrane to engage with CB₁ receptors, suppressing neurotransmitter release. The pharmacology of endocannabinoids is rapidly evolving with the discovery of new cannabinoid mimetics and novel CB receptor interactions that include orphan receptors and other CB receptors being proposed^[23-26]. All these receptor functions are either sensitive to, or are regulated by CB receptors and culminate in neuropathic pain and inflammation, suggesting that novel drug targets based around these proteins might be useful in treating neuroinflammation and neuropathic pain in different neurological conditions. In addition, activation of the CB receptor inhibits adenylate cyclase that then reduces the levels of the second messenger cyclic adenosine monophosphate (cAMP), thus regulating cellular mechanisms such as cell fate^[27]. CB receptor activation also inhibits voltage-dependent Ca²⁺ channels and activates inwardly rectifying K⁺ channels, a process that underlies CB-induced depression of excitatory neurotransmission^[28,29]. Thus, the CB system is a potentially

useful target for exploitation in neurodegenerative diseases such as MS.

AXONAL DAMAGE IN MS

Although MS is defined as an inflammatory demyelinating disease of the CNS, axonal loss is also a key feature of the disease. In MS patients, analysis of their spinal cord lesions suggested that the permanent loss in function was not primarily due to demyelination but due to axonal loss^[30]. Axonal loss is generally associated with inflammatory macrophage infiltration but is variable in MS lesions and can be severe in certain cases. For example, axonal density is reduced by 60%-70% in actively demyelinating lesions and is characterised by the presence of axonal spheroids, endbulbs, or focal accumulation of proteins^[31-33]. Although shadow plaques appear after injury that is consistent with an attempt to remyelinate axons, ongoing axonal injury is present^[34]. This suggests that during the early stages of remyelination axons are more susceptible to damage, a process that is related to the patterns of Na⁺ channels at the widened surfaces of the nodes of Ranvier^[35]. Axonal injury is also present in normal appearing white matter^[36,37] and may occur as a result of secondary Wallerian degeneration^[38]. However, this is not the only mechanism of axonal damage since two distinct patterns are observed: one that takes place within demyelinated lesions and correlates with lesional activity, while the other is diffuse axonal injury that is associated with inflammation and can additionally affect non-demyelinated nerve fibres^[33].

Axonal injury in MS is also selective to the size of axon fibres. Small calibre axons are more prone to injury compared to thick axons and may relate to thin axons requiring a higher energy demand in terms of their critical mass of mitochondria^[39]. Like MS, widespread axonal damage is also seen in experimental allergic encephalomyelitis (EAE)^[34,40]. However, experimental models of MS reveal different mechanisms of axonal injury and there is currently no agreement for which mechanism is relevant to MS patients. Axonal injury mechanisms may involve T-lymphocytes and antibodies as part of the adaptive immune response as well as components of the innate immune system, driven by macrophages and microglia. For example, axonal injury can be driven by an antigen-specific cytotoxic T cell response, induced in neurons and glia by the expression of pro-inflammatory cytokines, leading to neuronal death and axonal transection^[41-43].

Another mechanism of axonal damage relies on the production of auto-antibodies against cell surface molecules in neurons and axons. A subset of MS patients were described that mounted an antibody response against neurofascin, which is expressed on axons and oligodendrocyte processes at the nodes of Ranvier^[44]. Systemic injection of neurofascin during EAE exacerbates the clinical symptoms of the disease with severe levels of axonal injury within lesions^[44]. These observations suggest that auto-antibodies can directly mediate axonal damage. In MS lesions, axon damage is closely linked to

the presence of macrophages and microglia that are in intimate contact with axons^[3,5] and are known to produce a number of cytotoxic molecules including reactive oxygen and nitric oxide intermediates^[33]. The expression of these molecules, especially inducible nitric oxide synthase from macrophages, correlates with areas of axonal injury in acute EAE^[45]. In summary, axonal damage is a feature of MS and EAE and correlates with functional deficits. EAE models demonstrate that axonal damage occurs through a variety of mechanisms. However, recent reports have highlighted the role of mitochondrial injury and subsequent energy failure, induced by oxygen and nitric oxide free radicals as a possible mechanism of axonal injury. Understanding of the pathway to axon injury will aid in the discovery of new molecules for therapeutic intervention.

CANNABINOID IN MS

There is an abundance of evidence suggesting that MS patients gain symptomatic relief from cannabis extracts^[46]. For example, Sativex, an oral-mucosal spray containing Δ^9 -THC and CBD is anti-spasmodic and analgesic in MS patients^[47], while neuropathic pain associated with MS is relieved by dronabinol, an oral preparation of Δ^9 -THC analog^[15]. Meta-analysis has also revealed that CB-based preparations are superior in the treatment of MS-related neuropathic pain than the placebo, confirming their beneficial effects in symptomatic relief^[48]. This is consistent with the animal model of MS, EAE where treatment with exogenous cannabinoids controlled spasticity in chronic relapsing EAE models^[49].

Modulation of the endocannabinoid system is also apparent in MS and EAE such that brain levels of CB receptors are downregulated in EAE while plasma levels of endocannabinoids are increased in MS patients^[50,51]. Synthetic CB, HU-211, reduced the clinical severity of acute EAE in female Lewis rats as well as reducing inflammatory cell infiltration into the CNS^[52], while the WIN55, 212-2, a CB₁ and CB₂ receptor agonist reduces T cell differentiation and hence reduces EAE severity in Theiler's murine encephalomyelitis virus-induced demyelinating disease, a mouse model of chronic-progressive MS^[53], suggesting a key involvement of the CB receptors in the pathogenesis of EAE. Induction of EAE in CB₁ receptor-deficient mice causes rapid and progressively more neurodegeneration than in wild-type counterparts^[54].

In our highly cited study on the role of cannabinoids in EAE, reported that the cannabinoid system was neuroprotective during EAE since CB₁-deficient mice poorly tolerated inflammatory and excitotoxic insults and showed significant accumulation of neurodegeneration after EAE^[55]. We induced chronic relapsing EAE (CREAE) in wild type ABH, CB₁ gene (*Cnr1*)-deficient and congenic ABH. *Cnr1*^{+/+} mice with mouse spinal cord homogenate emulsified in complete Freund's adjuvant on days 0 and 7 and monitored clinical disease progression over time. Mice developed characteristic paralytic disease episodes followed by remission with an increas-

ing amount of residual deficit^[49,56]. Whilst disease induction in both CB₁-deficient and CB₁ wild-type mice were similar, CB₁-deficient mice exhibited significantly higher levels of residual deficit. This deficit was quantitated in an open-field activity chamber and confirmed that CB₁-deficient mice displayed significantly more immobility and paresis than wild type mice, accumulating significantly more axonal damage after relapses. CB₁-deficient mice also developed spasticity after only a single attack, which is not seen in wild type mice until after three to four relapses of disease^[49].

Numerous mechanisms cause neuronal death and axonal damage in EAE, including the influx of toxic ions such as Ca²⁺ and caspase-3-mediated apoptosis^[57]. CB₁-deficient mice demonstrated significantly lower levels of active caspase-3 during acute EAE compared with wild type mice, while caspase-3 was detected in dying axons, consistent with that observed in MS^[3]. These results suggested that the elevated neurodegeneration in CB₁-deficient mice may be due to caspase-3-mediated apoptosis and axonal damage and hence agonism of the CB₁ receptor pathway is neuroprotective and may control neurological symptoms such as tremor and spasticity^[49].

We also investigated whether glutamate toxicity can be regulated by cannabinoids^[55]. Glutamate excitotoxicity causes neuronal damage in both MS and EAE. For example, the glutamate antagonist, amantadine, reduces the relapse rate in MS patients^[58], while elevated glutamate levels have been observed in cerebrospinal fluid from MS patients^[59]. In EAE, the enhanced levels of glutamate agonists may result in aberrant astrocyte function, since activated astrocytes normally regulate glutamate levels through enzymes such as glutamate dehydrogenase and glutamine synthetase, both of which are down-regulated during EAE^[60,61]. The amount of CNS glutamate is also affected by abnormal changes in neuronal and glial glutamate transporters, all of which raise the levels of glutamate in the CNS during EAE and ultimately lead to the synthesis of mediators responsible for neuronal dysfunction^[59,62-64]. We reported that after *in vitro* stimulation of N-methyl-D-aspartic (NMDA) receptors in cerebellar granule cells, there was significantly more neuronal Ca²⁺ influx in CB₁-deficient mice than in wild type controls suggesting that the cannabinoid receptors may tonically regulate Ca²⁺ influx. The NMDA receptor antagonist MK-801 took longer to reduce Ca²⁺ back to basal levels in CB₁-deficient mice than in congenic controls while CB₁ agonism with CP55, 940 inhibited NMDA-induced Ca²⁺ influx in wild type mice but had no effect on CB₁-deficient mice. These results suggest that Ca²⁺ is not only dysregulated in the absence of CB₁ receptors but that postsynaptic control of NMDA-receptor activation is also compromised. These *in vitro* results were confirmed by *in vivo* injections of kainic acid, a specific agonist of the kainate receptor that mimics the effects of glutamate, since injection of kainic acid in CB₁-deficient mice induced seizures and mortality within 10 min, while no effect of kainic acid was observed in wild type or congenic wild type control mice despite using 50-fold higher doses.

Therefore, CB₁-receptors clearly regulate ionotropic glutamate receptor activity, leading to enhanced susceptibility to excitotoxic damage in CB₁-deficient mice.

Furthermore, we reported that CB₁ agonists protected mice against the consequences of CNS inflammation in models of experimental allergic uveitis (EAU)^[55]. EAU is an inflammatory disease of the eye and after sensitization with for example, interphotoreceptor retinal binding peptide in B10. RIII mice, the neuroretina is completely destroyed within 14–16 d^[65]. CB₁ receptor agonism with R(+)-WIN-55,212-2 and Δ^9 -THC both significantly inhibited photoreceptor damage without affecting inflammatory infiltrates suggesting that agonism of CB₁ is neuroprotective. Taken together, the results of our study demonstrated that cannabinoids protect against the neurodegenerative events in models of MS and EAU.

CANNABINOID-1 RECEPTOR-MEDIATED NEUROPROTECTION IN MODELS OF MS

A feature of MS is neuronal loss, and in MS patients, loss of spinal cord axons correlate with neurological disability together with reduced N-acetyl aspartate levels in chronic MS patients^[32]. Although axonal loss occurs early during the progression of MS^[66], once a threshold of 15%–35% axonal loss in mice has been reached, permanent disability results^[67,68]. In CB₁-deficient mice, significant axonal loss is evident after a single acute attack of EAE, suggesting that the mere presence of CB₁ is itself neuroprotective^[55]. In EAE and MS, the presence of axonal damage correlates with inflammation that produces a range of neurotoxic agents such as glutamate, cytokines as well as creating oxidative stress in the environment of the CNS and damaging the blood-brain barrier^[3,5,52,69-72].

Cannabinoids, however, can protect against acute hypoxia, excitotoxicity, oxidative and traumatic insults both *in vitro* and *in vivo*^[73-76]. Cannabinoids can also inhibit both pre- and post-synaptic glutamate induced calcium responses and thus inhibit neurotoxicity^[21,55,76], an effect that we showed to be CB₁-dependent^[55]. In accord with our observations, CB₁-deficient mice were more susceptible to NMDA and α -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA)/kainite glutamate receptor excitotoxicity^[55], however, Δ^9 -THC and CBD protected against NMDA-, AMPA- and kainite-agonist-induced cell death^[77,78]. Delta-9-THC also protected retinal neurons from death induced by peroxynitrite-mediated NMDA-induced toxicity^[79]. These observations demonstrate a clear neuroprotective role of cannabinoids in MS.

CANNABINOID-2-MEDIATED EFFECTS IN MODELS OF MS

Although much of the focus in MS is devoted to CB₁ receptor pharmacology, the recognition that CB₂ receptors possess immunomodulatory properties has led to an increased focus on CB₂ receptors as potential thera-

peutic targets. Unlike CB₁ receptors, activation of the CB₂ receptor is not psychoactive and therefore targeting CB₂ receptors with selective agonists is a promising therapeutic avenue that is immunomodulatory without being psychoactive. For example, early indications came from experiments that showed that the administration of WIN55212-2, which functions as a CB₁ and CB₂ receptor agonist, attenuated the progression of EAE in C57BL/6 mice immunized with myelin oligodendrocyte-derived glycoprotein35-55 (MOG35-55)^[80]. Furthermore, a selective antagonist of the CB₁ did not modulate the protective effect of WIN55212-2 but a selective antagonist of the CB₂ receptor blocked the effects of WIN55212-2. This led to the suggestion that the protective effects of WIN55212-2 was mediated through the CB₂ receptor^[80]. However, later studies showed that the CB₁ receptor might play a neuroprotective role in the latter stages of EAE^[49,54].

Other highly selective CB₂ receptor agonists such as O-1966 are also useful in the fight against MS since they do not produce psychoactive effects, determined by its low affinity to CB₁ receptors^[81]. Administration of O-1966 in a chronic (C57BL/6/MOG), relapsing-remitting and an adoptive transfer model attenuated disease progression and improved motor function^[82]. In addition, encephalitogenic T cells derived from CB₂-deficient mice were shown to be more aggressive in terms of CNS infiltration and increased severity of EAE, despite displaying similar levels of proliferation, apoptosis and cytokine production in the spleen as wild-type T cells^[83]. Moreover, treatment of mice with CB₂ receptor agonists attenuate white cell trafficking across the blood-brain barrier while dendritic cells differentiated in the presence of selective CB₂ receptor agonist and inhibited T cell proliferation and shifted cytokine responses from inflammatory to anti-inflammatory molecules^[82]. Recently, it has been shown that high concentrations of IFN- γ disrupts P2X₇ purinergic receptor signalling and thus inhibit the neuroprotective effects of endogenous cannabinoids^[84]. Therefore, inhibiting IFN- γ by exogenous CB₂ agonists represents an obvious therapy to prevent the disruption of P2X₇ signalling.

VALUE OF ANIMAL MODELS IN THE STUDY OF MULTIPLE SCLEROSIS

The validity of EAE as a model of MS is a topic of active debate with some researchers contenting that it is unsuitable due to its inability to mimic some of the pathological, immunologic and chronic features of MS. For example, EAE is usually monophasic whereas MS displays chronic relapsing features. Histological and magnetic resonance imaging data demonstrate axonal and cortical damage in MS but not in some models of EAE^[85]. The extravasation of red blood cells into the CNS of swiss jim lambert mice and Lewis rats with EAE are not typical of MS^[86,87], while encephalitogenic regions associated with myelin basic protein (MBP) or proteolipid protein normally activate more CD4⁺ than CD8⁺ T cells

but in inflammatory MS, CD8⁺ T cell predominate^[88-90]. However, there are many other immunological differences between mouse and human MS and these must be overcome if greater levels of success in drug development for human MS are to be achieved^[91].

Several other points are worth considering in terms of the use of EAE as an animal model: (1) EAE provides little insight into the progression of MS in terms of the small amounts of demyelinated axons in EAE compared to MS while mice with EAE rarely exhibit ongoing functional deterioration that MS patients often display^[92,93]; (2) The use of C57BL/6 mice in EAE studies limits the investigation of the mechanism of relapsing-remitting forms that more commonly affect MS patients (Ransohoff *et al*^[93], 2002); (3) Treatment with factors that exert neurobiological effects also impact on immune and inflammatory cells and thus making results difficult to interpret (Ransohoff *et al*^[93], 2002); and (4) EAE is generally generated using antigens that affect CD4⁺ T cells while CD8⁺ T cells that predominate in MS lesions are overlooked^[92,94]. Likewise, the role of B cells is largely neglected despite recent data that demonstrate their importance in the pathogenesis of MS^[95,96].

In summary, EAE has a long history in the fight against MS, however, its predictive value for treatment efficacy is poor. Nevertheless, it is widely used as a first-line animal model of MS and has provided mechanistic insights into the neuroinflammatory aspects of MS.

LIMITATIONS OF ANIMAL MODELS OF MS

One of the biggest limitations on the use of EAE as a model of MS is the fact that disease has to be induced with complete Freund's adjuvant and heat-inactivated *Mycobacterium tuberculosis* rather than mimicking a spontaneous disease like MS. This leaves very little room for disease pathways and fails to represent the complexity of disease inducing mechanisms in MS. Demyelination, a feature of MS is also not obvious in all EAE models while the time course for disease manifestation may be days, EAE more closely resembles post-infectious acute demyelinating events^[97]. In contrast, MS develops over years with patients presenting with more protracted epitope spreading than that observed in EAE mice^[98]. There are also many other immunological differences between EAE and MS that need consideration in the development of potential therapies for MS.

Therapeutic developments from EAE have translated poorly to human MS. For example, only a few molecules that showed efficacy in EAE have been successful in MS trials. One of these molecules is Glatiramer acetate, a synthetic amino acid copolymer originally designed to mimic encephalitogenic MBP, but instead suppresses EAE by other mechanisms and reduced MS relapses by 30%^[99,100]. However, the efficacy of Glatiramer acetate has been questioned by a systematic Cochrane review which calls into question the use of Glatiramer acetate in

MS^[101]. Tysabri (Natalizumab) and Gilenya (fingolimod) are the only two other drugs that have been licence for use in human MS. Natalizumab binds to the $\alpha 4 \beta 1$ and $\alpha 4 \beta 7$ integrins and blocks binding to their endothelial receptors (VCAP-1 and mucosal addressin-cell adhesion molecule 1, thereby attenuating inflammation and ongoing inflammation^[102,103]. Fingolimod is a sphingosine-1-phosphate-receptor modulator that prevents egress of lymphocytes from lymph nodes and significantly improved relapse rates compared to placebo controls^[104]. At present therefore, it remains unclear why pre-clinical EAE studies predict treatment efficacy in human MS so poorly, however, EAE is still an important first-line model system in the development of new treatments for MS.

BETTER ANIMAL MODELS OF MULTIPLE SCLEROSIS

To make greater progress, better animals models of MS need to be considered when testing new drugs. One fact that to be borne in mind is the fact that MS is a heterogeneous disease in terms of genetics, environmental effects, disease course, pathological treatments and treatment responsiveness^[105]. Currently the majority of experiments are performed in inbred strains while clinically, the molecular mechanisms that determine the efficacy during prolonged follow-up of hundreds of patients are far more complex. Thus, it is likely that genetic differences account for some of the inter-patient differences in clinical efficacy of various drugs. Improved animal models may take into account the therapeutic effect of drugs in more than one animal model, treatments to be instigated after disease onset, long-term disease periods, use spontaneous disease models and incorporate human risk. The use of partially humanized mouse models to address the genetic and disease variability are a step in the right direction towards developing better therapeutics for MS^[106].

SUMMARY AND FUTURE PROSPECTS

Studies have shown that antagonism of the CB₁ receptor pathway contributes to neuronal damage in CREAE and the relative worsening of tremor and spasticity. The addition of exogenous CB₁ agonists derived from cannabis afforded significant neuroprotection from the consequences of inflammatory CNS disease in EAE and EAU models. Although clear neuroprotective benefits of cannabinoids have been demonstrated, the unwanted psychotropic effects need to be addressed. The adverse psychotropic effects, its role in appetite, pain and cognition together with the observation that the CB₁ receptor is downregulated in some neurodegenerative diseases limits the usefulness of cannabinoids in the treatment of MS. Alternative treatments may be more useful and includes the development of drugs based on CBD, the non-psychoactive part of cannabis that possess anti-inflammatory and anti-oxidant properties. Furthermore,

the CB₂ receptor is being recognised as a potential target since it regulates neuroinflammation and neurogenesis while being non-psychoactive.

In summary, the endocannabinoid system exerts multiple actions and may be useful in the development of therapies to treat neurodegenerative diseases. However, the biggest challenge remains, namely the development of drugs that lack the adverse psychoactive side effects.

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Secondary prevention of ischaemic stroke

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Abstract

In spite of a documented reduction in incidence in high-income countries over the last decades, stroke is still a leading cause of death and disability worldwide. With the ageing of the population stroke-related economic burden is expected to increase, because of residual disability and its complications, such as cognitive impairment, high risk of falls and fractures, depression and epilepsy. Furthermore, because of the substantial rate of early and long-term vascular recurrences after the first event, secondary prevention after cerebral ischaemia is a crucial issue. This is even more important after minor stroke and transient ischaemic attack (TIA), in order to reduce the risk of potentially more severe and disabling events. To accomplish this aim, acute long-term medical and surgical treatments as well as

lifestyle modifications are strongly recommended. However, apart from the well-established indications to thrombolysis, studies in acute phase after a first stroke or TIA are scarce and evidence is lacking. More trials are available for long-term secondary prevention with different classes of drugs, including antithrombotic medications for ischaemic events of arterial and cardiac origin, especially related to atrial fibrillation (antiplatelets and anticoagulants, respectively), lipid lowering agents (mainly statins), blood pressure lowering drugs, surgical and endovascular revascularization procedures.

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Key words: Stroke; Transient ischaemic attack; Secondary prevention; Antiplatelets; Anticoagulants; Medical stroke treatment; Carotid stenosis

Core tip: Aggressive and combination treatments in the acute phase after transient ischaemic attack or minor stroke have been shown to be beneficial in few studies, but results of ongoing randomized trials are required. On the other side, in long-term prevention the most important innovation is the advent of new anticoagulant agents for stroke prevention in atrial fibrillation. Recent trials showed efficacy and safety of thrombin and factor Xa inhibitors, compared to vitamin K antagonists, whose use is hampered by several limitations. These new drugs will potentially increase the number of patients treated according to guidelines, thus preventing a remarkable proportion of strokes.

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INTRODUCTION

Stroke is a leading cause of death and disability worldwide^[1] and its burden on global health-care expenditure is supposed to increase, mainly because of the ageing population. Incidence of stroke is higher than that of myocardial infarction in some regions^[2] and in spite of a mild reduction of incidence in developed countries in the last four decades, a rise of cerebrovascular events has been observed in low-income countries^[3]. Stroke has also substantial indirect costs related to complications, such as post-stroke dementia, depression, falls, fractures and epilepsy.

Almost 90% of all cerebrovascular events depends on few medical conditions, including bad habits (*e.g.*, smoking, physical inactivity, diet, alcohol excess, stress) as well as arterial hypertension, cardiac diseases, diabetes, and high ratio of apolipoprotein B to apolipoprotein A1^[4]. Lifestyle modifications as well as medical and surgical preventive strategies may, therefore, reduce the risk of stroke. Although primary prevention remains a milestone, effective secondary prevention plays also a fundamental role in this regard. About 30% of strokes occur in individuals with a previous transient ischaemic attack (TIA) or stroke^[2], and more than 50% occur in subjects with previous vascular events of any kind^[5]. Much progress has been made in the last decades in stroke prevention, and there is now evidence from randomized trials that several treatments are effective in preventing recurrences (Figure 1). In particular, at least 80% of recurrent events might be prevented with the use of a comprehensive approach that includes dietary modifications, physical exercise, blood-pressure lowering drugs, antiplatelet therapy, and statins^[6].

The effect of all these different interventions on stroke risk can only be derived from population-based studies. In this regard, it has been shown that the risk of recurrent stroke in all patients in a United Kingdom population-based study after a first-ever TIA or non-disabling stroke in the period 1981-1986 was higher than the risk of the same population during 2002-2010^[5]. These findings were corroborated by a recent review^[7] of randomized controlled trials in secondary prevention after stroke and vascular events. Recurrent stroke and vascular events rates declined substantially in the last 5 decades, mostly because of improved blood pressure (BP) control and more frequent use of antiplatelet therapy.

In this review we will focus on medical treatment for secondary prevention in patients with first-ever TIA or ischaemic stroke and separate acute secondary prevention (*i.e.*, prevention in the first hours or days) from long-term secondary prevention. We will also deal with surgical procedures and endovascular techniques for prevention of recurrences in carotid stenosis as well as therapeutic strategies in stroke associated with patent foramen ovale (PFO).

EARLY SECONDARY PREVENTION

Background

The risk of recurrent stroke is higher during the first 90 d

after an index event, longitudinal studies showing that approximately 1 out of 2 recurrences in the first year post-stroke occurs within the first 90 d^[8-12]. The 90-d risk of recurrence after a TIA has been reported to be as high as 17%, with the greatest risk in the first week^[13,14]. In this regard, prospective studies have shown that stroke risk after a TIA or minor stroke is 3.1% (95%CI: 2.0-4.1) at 2 d and 5.2% (3.9-6.5) at 7 d^[7,15]. After a stroke or TIA, the risk of recurrence is still high in the first month (4%) and year (12%) but falls down to about 5% per year thereafter^[15]. For this reason, secondary prevention should be started urgently in the acute phase after TIA or minor stroke. Identifying patients at higher risk of recurrence is, therefore, crucial. For such a purpose several risk scores have been created. Comparison between these scores is hampered by differences in index event, methodology and factors taken into account. To date, the ABCD2 score combined with diffusion-weighted imaging (DWI) data (ABCD2I)^[16-19] appears as the best predictor of the early risk of stroke after a TIA. The recently validated ABCD3I score, including carotid stenosis in addition to abnormal DWI, showed even higher predictive power, although c-statistics were less convincing in the validation set^[20].

The evidence for acute treatment after TIA and minor stroke is derived from few small randomized trials, extrapolation of results from larger trials in other settings, and on two non-randomized studies of a combination of preventive treatment started urgently^[21]. The Early Use of Existing Preventive Strategies for Stroke (EXPRESS) study^[22] was nested in a population-based study of all acute TIAs and strokes in the Oxfordshire and compared urgent assessment and treatment (phase 1) with appointment-based clinic assessment and primary care-initiated treatment (phase 2). Results demonstrated a reduction of the 90-d risk of stroke recurrence from 10.3% in phase 1 to 2.1% in phase 2. Similarly, in the SOS-TIA study^[23], urgent intensive treatment was associated with a better prognosis and a low rate of recurrent events.

Based on the results of these studies, guidelines now recommend that patients with suspected TIA should be quickly referred to a TIA clinic or to a medical centre providing rapid evaluation and appropriate treatment. In such patients, urgent vascular imaging (ultrasound, cranial tomography angiography or magnetic resonance angiography) in addition to standardized emergency diagnostic tests, including cranial tomography or magnetic resonance imaging, electrocardiography and laboratory tests are recommended^[24,25].

However, the relative contribution of antithrombotic therapy, revascularization techniques, statins and antihypertensive drugs in stroke prevention remains uncertain.

Antithrombotic treatment

Aspirin is the only antiplatelet drug evaluated in the acute phase of stroke. Given within 48 h of onset of major ischaemic stroke it reduces 14-d morbidity and mortality, mostly by reducing the risk of early recurrent stroke^[26]. However, the optimal dose of aspirin has not been stud-

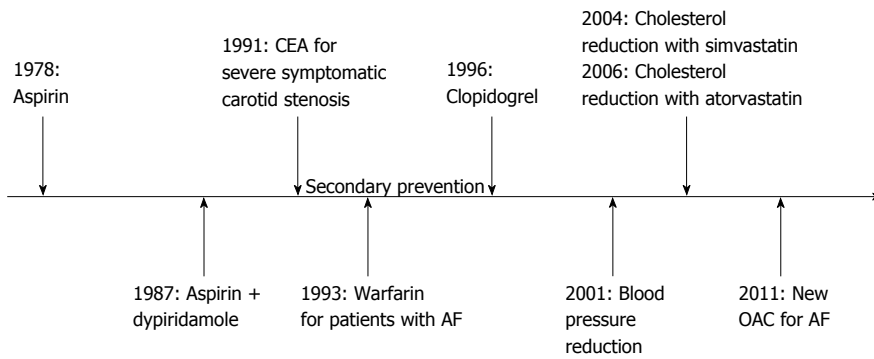


Figure 1 Medical and surgical treatments which have been proved to be effective for secondary prevention after stroke or transient ischaemic attacks in the last decades. OAC: Oral anticoagulants; CEA: Carotid endarterectomy; AF: Atrial fibrillation.

ied in head-to-head comparison. A recent meta-analysis by Chen *et al.*^[27], including studies in which daily doses of aspirin ranged between 160 to 325 mg, found no heterogeneity of effect on different outcomes. Similarly, optimal timing of initiation of aspirin therapy has not been studied in randomized trials. Though unproven, it seems reasonable to start aspirin therapy as soon as possible, ideally within 48 h of stroke onset. Furthermore, aspirin effect may be negligible in patients at high risk for early recurrence, such as those with stroke or TIA due to arterial thromboembolism^[28]. Alternative approaches in these cases, based on the combination of multiple antiplatelet agents with different mechanisms of action, seem, therefore, reasonable. The combination of aspirin and clopidogrel is, actually, more effective than aspirin alone in preventing vascular recurrences and death in acute coronary syndromes^[29]. In line with these findings, there is some evidence that a short course of more intensive antiplatelet treatment might be effective also in the acute phase after TIA and minor stroke^[21]. Among 731 patients with acute ischaemic stroke or TIA (onset within 3 h) enrolled in five small trials, the addition of clopidogrel to aspirin was associated with a nonsignificant trend toward a reduction in recurrent stroke and an increase in major bleedings^[30]. Although dual therapy seems not effective in lacunar strokes^[31], it may be more effective than single antiplatelet therapy in preventing early recurrence in patients with acute symptomatic atherothrombosis, according to the recent Clopidogrel in High-risk patients with Acute Nondisabling Cerebrovascular Event trial^[32] conducted on a Chinese stroke population. Because of these inconsistencies, the results of three ongoing trials investigating the safety and efficacy of dual and triple antiplatelet therapy^[33-35] are required before definite recommendations can be made.

Antihypertensive agents

An increase of BP values (even in previously normotensive patients) during the acute phase of ischaemic stroke, followed by a spontaneous decline starting within 90 min of symptoms onset^[36] is common finding after stroke. Although extreme arterial hypertension is clearly detrimental, as it might exacerbate oedema and favor the haemorrhagic transformation of ischemia, a moderate in-

crease of BP values during the acute phase of stroke may improve the perfusion of the ischaemic tissue, leading to better outcomes. Conversely, an impairment of cerebral perfusion by excessive BP lowering may exacerbate the ischaemic damage, an effect that is even more evident in conditions of chronically damaged autoregulation, such as in elderly subjects, or in cases of severe carotid stenosis^[37]. Unfortunately, the ideal range of BP values to be maintained early after stroke has not yet been determined. Larger trials with well-defined criteria are needed. At present, many clinicians in their daily clinical practice consider a BP greater than 220/120 mmHg an indication to active BP lowering^[36]. However, this threshold might be different in cases with minor cerebral damage, such as TIA or minor strokes, where the reduction of cerebral perfusion is likely to be of less concern. In line with this view, antihypertensive therapy started immediately after the acute event did not negatively influence the outcome in the EXPRESS^[22] and SOS-TIA^[23] studies.

Although antihypertensive medications are effective in the long-term secondary prevention after stroke and TIA^[38], the issue of early BP lowering after a cerebrovascular event is still uncertain, recent trials suggesting no benefit^[39-41]. In the Controlling Hypertension and Hypotension Immediately Post-Stroke trial^[39], oral nimodipine started within 48 h after ischaemic stroke onset in 350 patients was associated with a higher mortality rate and similar functional outcome. The Scandinavian Candesartan Acute Stroke Trial^[41], comparing candesartan to placebo given within 30 h of ischaemic or haemorrhagic stroke, showed no difference in the composite endpoint of vascular death, myocardial infarction or stroke after 6-mo follow-up. Similarly, in the Continue or Stop Post-Stroke Antihypertensive Collaborative Study trial^[40], continuation of antihypertensive therapy during hospitalization for ischaemic stroke was not associated with a reduction of 6-mo mortality or cardiovascular event rate, when compared to discontinuation of preexisting antihypertensive drugs.

Lipid lowering agents

In the acute phase after a cerebrovascular event statins might be beneficial because of their pleiotropic effects. These include a neuroprotective effect, limiting damage

and improving recovery (in particular for major strokes), as well as an antithrombotic effect, which in turns may prevent early recurrences (especially in TIAs and non disabling strokes). The benefits related to neuroprotective effects are perhaps more evident in the first 3 to 7 d after stroke and with higher doses of statins, as suggested by some observational studies and clinical data^[42-44]. However, data from the fast assessment of stroke and transient ischaemic attack to prevent early recurrence pilot trial are discordant^[45]. In this recent trial patients who presented with TIA or minor stroke within 24 h of symptoms onset were randomly assigned to simvastatin or placebo. No evidence of benefit of simvastatin was shown in the primary outcome of recurrent strokes within 90 d. Furthermore, a recent meta-analysis, including 8 randomized controlled trials, found insufficient evidence to establish safety and effectiveness (also in terms of secondary prevention) of statin use in the acute phase after ischaemic stroke^[46]. Similarly, initiation of statin therapy within 14 d following the onset of acute coronary syndromes does not reduce significantly the risk of death, myocardial infarction or stroke up to 4 mo^[47]. The major ongoing studies aimed at further assessing the benefit of early statin administration mainly focus on functional outcomes instead on the prevention of early recurrences^[46].

Glycemic control

Hyperglycemia is common during acute ischaemic stroke, being present in more than 40% of patients, mostly among those with a history of diabetes^[48,49], and it is associated with worse outcome. So far, only 1 randomized efficacy trial of hyperglycemia treatment in acute stroke, the Glucose-Insulin-Stroke Trial-UK^[50], has been reported. Nine hundred and thirty-three patients with acute ischaemic stroke within 24 h of symptom onset, not previously treated with insulin, were randomized to unblinded intravenous treatment with insulin, potassium, and glucose versus saline for 24 h. Although the results of this trial in terms of clinical outcomes were neutral, the key questions remain unanswered because of the trial design. Thus, the definitive efficacy and safety of earlier and greater reductions in serum glucose levels during acute ischaemic stroke remain to be studied. In the meantime, it is reasonable to treat hyperglycemia to maintain blood glucose levels in a range of 140 to 180 mg/dL, preventing hypoglycemia with close monitoring, according to the current American Diabetes Association guidelines for glycemic control^[51]. No data are available for the treatment of diabetes in the setting of TIA or minor strokes, but it seems reasonable to apply the same recommendation as in major strokes. Furthermore, information concerning the value of hyperglycemic control in early secondary prevention of stroke, as well as in long-term secondary prevention, is lacking.

LONG-TERM SECONDARY PREVENTION

Evidence from randomized controlled trials for long-term secondary prevention of ischaemic stroke is more

robust than for acute treatment.

Antithrombotic treatment

Ischaemic stroke is a heterogeneous clinical condition and the end-result of different underlying mechanisms: atherothrombosis (30%), cardioembolism (20%), small-vessel disease (25%). Approximately 25% of cases have no detectable cause (cryptogenic). Elucidating the mechanism of stroke is crucial to select the most appropriate therapeutic strategy and prevent further events.

Cerebral ischaemia of arterial origin: Antiplatelet agents are the cornerstone for secondary prevention of ischaemic events in patients with noncardioembolic strokes. Commonly used antiplatelet agents are aspirin, dipyridamole, and clopidogrel. Aspirin is an irreversible inhibitor of cyclooxygenase-1, which in turn inhibits the formation of thromboxane A2. Dipyridamole increases cyclic AMP by inhibiting platelet phosphodiesterase E5. Clopidogrel is a thienopyridine P2Y₁₂ adenosine diphosphate receptor blocker. Ticlopidine, which has the same mechanism of action as clopidogrel, is no longer recommended because of its side effects.

Aspirin, the most commonly used antiplatelet drug, is recommended for secondary prevention after cerebral ischaemia of arterial origin at a dose of 30-325 mg daily^[24,25,52]. In order to reduce bleeding complications, after the acute phase (generally 1 or 2 wk after the event) the aspirin dose may be reduced to 75-100 mg^[52]. However, in a meta-analysis of trials for secondary prevention of stroke, the relative reduction in stroke risk achieved by aspirin is small, being only 13%^[53]. To date, the three large trials^[54-56] which investigated whether vitamin K antagonists (VKAs) are more effective than aspirin for this indication, showed no difference, irrespective of the intensity of anticoagulation. Several trials have studied other antiplatelets as an alternative or in addition to aspirin. The second European Stroke Prevention Study (ESPS2) trial^[57] and the European/Australasian Stroke Prevention in Reversible Ischaemia Trial (ESPRIT)^[56] investigated the efficacy of the combination of aspirin plus dipyridamole versus aspirin alone. Meta-analysis showed a relative risk reduction of 18% in the vascular events in favour of the combination therapy^[58]. In the Clopidogrel versus Aspirin in Patients at Risk of Ischaemic Events (CAPRIE) study, clopidogrel achieved a non-significant 7.3% reduction in relative risk of the composite outcome of stroke, myocardial infarction or vascular death, when compared to aspirin^[59]. The combination of aspirin plus clopidogrel was compared with clopidogrel alone in the Aspirin and Clopidogrel compared with Clopidogrel Alone after Recent Ischaemic Stroke or TIA in High-risk Patients (MATCH) trial^[60] and with aspirin monotherapy in the Clopidogrel for High Atherothrombotic Risk and Ischaemic Stabilization, Management, and Avoidance (CHARISMA) trial^[61]. Both trials showed higher rates of bleeding complications in the dual antiplatelet arm which overcome any beneficial effect. A more recent study, comparing clopidogrel plus aspirin with aspirin

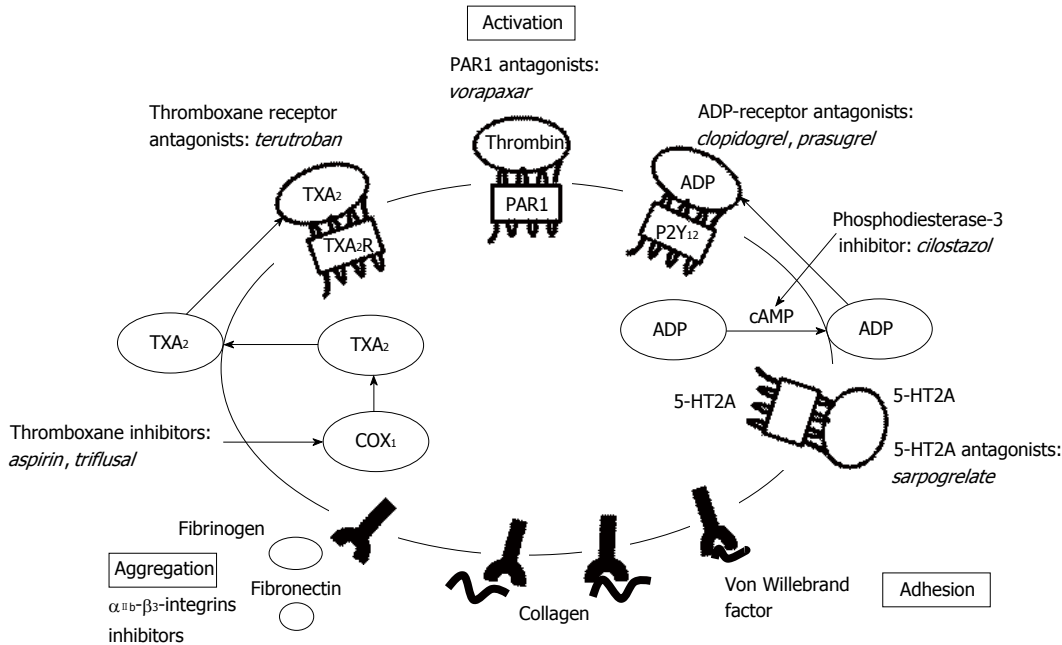


Figure 2 Old and new antiplatelet agents and their mechanism of action. PAR1: Proteinase activated receptor 1; ADP: Adenosine diphosphate; TXA₂: Thromboxane A₂; cAMP: Cyclic adenosine monophosphate; COX-1: Cyclooxygenase-1; 5-HT_{2A}: 5-hydroxytryptamine (serotonin) receptor 2A.

alone in recent lacunar strokes confirmed these results^[31]. Although in indirect comparisons the combination of aspirin plus dipyridamole was more effective than clopidogrel^[62], the non-inferiority Prevention Regimen for Effectively Avoiding Second Strokes (PROFESS) trial did not confirm this result, showing a similar rate of recurrent stroke in both treatment groups^[63]. Furthermore, the rates of discontinuation and of major systemic hemorrhages were higher in the aspirin plus dipyridamole group.

Current therapies have several limitations, including a weak inhibition of platelet function with modest clinical effect, blockade of only one platelet pathway, slow onset of action (*i.e.*, for clopidogrel), high bleeding risk, and interpatient response variability with poor inhibition of platelet activity. The concept of “antiplatelet resistance” has been proposed for those patients experiencing recurrences during correct antiplatelet therapy. It occurs when the drug biochemically fails to inhibit platelet activation, as measured by *in vitro* platelet function assays. Proposed mechanisms responsible for this phenomenon include drug-drug resistance and genetic polymorphisms of antiplatelet metabolism or drug-receptor site. However, because of the lack of a standardized assay and the poor correlation between antiplatelet resistance and recurrent clinical events, currently there is no indication to screen patients for antiplatelet resistance^[64,65]. For clopidogrel, in particular, it seems prudent to avoid interactions of drugs with a well-known effect on hepatic cytochrome p450, as it has been demonstrated that these drugs might affect clopidogrel metabolism^[64]. Consequently, in the last years, other antiplatelet agents with different mechanisms of action have been investigated and compared to aspirin (Figure 2). Triflusal is a thromboxane inhibitor structurally related to aspirin with a similar efficacy in

preventing vascular events, but a lower risk of bleeding, in spite of more non-haemorrhagic adverse gastrointestinal events^[66]. Cilostazol is a phosphodiesterase-3 inhibitor which represents a promising alternative to aspirin in Asian people with stroke. A meta-analysis including two randomized clinical trials concluded that cilostazol is more effective than aspirin in the prevention of vascular events secondary to stroke, causes more frequently minor adverse effects, but less frequently bleeding complications^[67]. The Prevention of Cardiovascular Events in Ischaemic Stroke Patients with High Risk of Cerebral Haemorrhage (PICASSO) trial is an ongoing study aimed at randomizing patients with previous stroke and high bleeding risk (*i.e.*, patients with a symptomatic or asymptomatic old cerebral haemorrhage) to aspirin or cilostazol (and to probucol)^[68]. Sarpogrelate, a selective inhibitor of 5-hydroxytryptamine receptors, has been investigated in comparison with aspirin and turned out to be non-inferior to the latter for prevention of stroke recurrences, with fewer bleeding events^[69]. Terutroban, which is a specific antagonist of the thromboxane A₂ receptor, was no more effective than aspirin for prevention of stroke in the large Prevention of Cerebrovascular and Cardiovascular Events of Ischaemic Origin with Terutroban in Patients with a History of Ischaemic Stroke or TIA (PERFORM) trial^[70]. Finally, vorapaxar is a novel antiplatelet agent that selectively inhibits the cellular action of the thrombin through antagonism of proteinase activated receptor 1. In a recent trial^[71] in which vorapaxar was compared to placebo in patients with vascular events, it reduced the risk of cardiovascular death or ischaemic events in patients with stable atherosclerosis who were receiving standard therapy. However, it increased major bleedings, including intracranial haemorrhages.

On the basis of these trials, international guidelines

Table 1 Antiplatelet therapy after non-cardioembolic stroke or transient ischaemic attack

Options		Recommendation
AHA/ASA ^[25]	Aspirin (50-325 mg); aspirin plus dipyridamole; clopidogrel	Aspirin, aspirin plus dipyridamole or clopidogrel
ESO ^[24]	Aspirin; aspirin plus dipyridamole; clopidogrel; triflusal	Clopidogrel or aspirin plus dipyridamole
ACCP ^[52]	Aspirin (75-100 mg); clopidogrel; aspirin plus dipyridamole; cilostazol	Clopidogrel or aspirin plus dipyridamole

AHA/ASA: American Heart Association/ American Stroke Association; ESO: European Stroke Organisation; ACCP: American College of Chest Physicians.

Table 2 CHADS₂, CHA₂DS₂VASc, and HAS-BLED risk stratification methods

CHADS₂		
Congestive heart failure	1	
Hypertension	1	
Age ≥ 75 yr	1	
Diabetes mellitus	1	
Stroke, TIA, or thromboembolism	2	
Maximum score	6	
CHA₂DS₂VASc		
Congestive heart failure/LV dysfunction	1	
Hypertension	1	
Age ≥ 75 yr	2	
Diabetes mellitus	1	
Stroke, TIA, or thromboembolism	2	
Vascular disease (previous MI, PAD, or aortic plaque)	1	
Age 65-74 yr	1	
Sex category (female sex)	1	
Maximum score	9	
HAS-BLED		
Hypertension	1	
Abnormal renal/liver function	1 point each	
Stroke	1	
Bleeding history or predisposition	1	
Labile INR	1	
Elderly (age > 65 yr)	1	
Drugs (e.g., concomitant antiplatelet/NSAID) or alcohol	1 point each	
Maximum score	9	

TIA: Transient ischaemic attack; LV: Left ventricular; MI: Myocardial infarction; PAD: Peripheral artery disease; NSAID: Nonsteroidal inflammatory drugs; INR: International normalized ratio.

still vary, but most recommend aspirin plus dipyridamole or clopidogrel as first line therapy in long-term secondary prevention of stroke. If these are not available or contraindicated, aspirin monotherapy, triflusal and cilostazol are alternatives (Table 1). However, even if both aspirin plus dipyridamole and clopidogrel alone are statistically superior compared to aspirin alone, clinical superiority is at best minimal. Furthermore, if tolerance profile is included in judgment of efficiency, then aspirin or clopidogrel monotherapy should be advised as the first-line therapy, since aspirin plus dipyridamole and aspirin plus clopidogrel carry higher rates of discontinuation and of bleeding complications, respectively. If also economic considerations are taken into account, then aspirin therapy remains the best choice^[72].

Patients who present with a first or recurrent stroke are commonly already on antiplatelet therapy. To date, for patients experiencing a stroke during aspirin therapy, there is no evidence that increasing the dose of aspirin

provides additional benefit. Moreover, although alternative antiplatelet agents are often considered, no single agent or combination has been studied. In such a situation it is important to take into account the compliance to therapy, as well as other vascular risk factors and possible different stroke mechanisms.

Cerebral ischaemia caused by atrial fibrillation:

Roughly 20% of all TIAs and ischaemic strokes have a cardiac origin, most commonly caused by atrial fibrillation (AF). AF is associated with a five-fold increased risk of stroke, the yearly risk varying from 0.2% in patients with AF alone to more than 10% in individuals with other risk factors^[73]. The major risk factors for stroke in AF patients are a history of stroke or TIA, advancing age, hypertension, systolic BP (SBP) > 160 mmHg, and diabetes. These and other risk factors have been used to create several scores for stroke risk calculation, in order to recognize patients who could benefit from anticoagulation. The most widely used model for stratification of stroke risk is CHADS₂ score (Table 2). However, although the score is simple and has been externally validated it does not reliably discriminate between patients with low risk, who do not need anticoagulation, and patients at intermediate risk, who do^[73]. This has prompted the development and the external validation in independent cohorts of a new score, the CHA₂DS₂VASc score (Table 2). Because of the inclusion of newly recognized risk factors for stroke, this score allows a better stroke risk stratification in patients who are considered at low or intermediate risk for stroke^[74]. Beside quantification of stroke risk, another important issue is the stratification of bleeding risk associated with anticoagulation. Among the schemes created for such a stratification, the most frequently used is the HAS-BLED score (Table 2), which was originally derived from a cohort of 3456 patients with AF and a one-year follow-up^[75]. HAS-BLED score, which has been validated in external cohorts, correlates with the risk of intracranial bleeding, a score of more than 3 identifying patients at high bleeding risk. In these conditions, it does not necessarily mean that anticoagulation should not be initiated but that such therapy warrants special caution, control of modifiable risk factors, and close monitoring^[73]. Difficulty arises in clinical practice when patients have key risk factors for both ischaemic stroke and major bleeding (e.g., age, hypertension, or previous stroke). Among these factors, previous stroke and increasing age are more strongly associated with ischaemic stroke than with intracranial bleeding^[76,77].

Recommendations for patients with AF and a his-

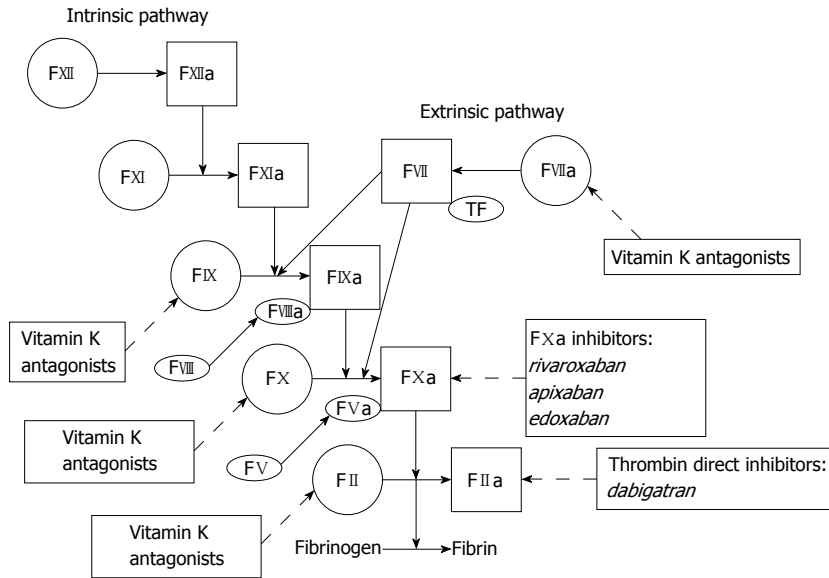


Figure 3 Vitamin K antagonist and new anticoagulant mechanism of action. F: Factor.

tory of stroke and TIA are based on the pooled effect of anticoagulation in primary and secondary prevention studies, as data on secondary prevention are limited to few studies. The first of these studies is the European AF Trial^[78], in which patients were randomly allocated to VKAs, 300 mg/d-aspirin, or placebo. VKAs turned out to be the most effective treatment, despite a higher number of major haemorrhages. The findings were confirmed in patients with previous TIA or ischaemic stroke enrolled in the Stroke Prevention in AF III trial^[79], which was halted early because patients assigned to fixed low-dose VKA (INR 1.2-1.5) plus aspirin 325 mg daily had four times more ischaemic events, compared to those allocated to regular VKA dose. A further trial in patients with recent cerebral ischaemia and AF found 15% reduction of recurrences with VKA than with indobufen^[80]. Adjusted-dose warfarin (target INR 2.0-3.0) turned out to be significantly more effective than the combination of aspirin plus clopidogrel in the AF Clopidogrel Trials with Irbesartan for prevention of Vascular Events (ACTIVE)^[81,82]. ACTIVE W was stopped prematurely because of an excess of primary outcomes in the aspirin plus clopidogrel group than in warfarin group. ACTIVE A included patients who were not eligible for VKAs and showed a small advantage for the combination of clopidogrel and aspirin compared to aspirin alone in preventing ischaemic events, but with more bleedings.

Despite their effectiveness, VKAs have many limitations, including slow onset and offset of action, interindividual variability in their effect, narrow therapeutic index, food and drug interactions and reduced synthesis of all vitamin K-dependent proteins^[83]. It has been calculated that at least a third of patients with AF who are at risk for stroke are either not started on VKAs therapy or discontinue the therapy. These limitations have prompted to the development and assessment of other agents that target the coagulation cascade at sites different from those

of classic VKAs: factors Xa and direct thrombin inhibitors (Figure 3). The first of these agents, Ximelagatran, a factor Xa inhibitor, was investigated in two trials^[84,85]. It resulted as effective as warfarin in prevention of stroke with fewer bleedings, but it turned out to be hepatotoxic and was then withdrawn. Dabigatran, a thrombin inhibitor, was compared to dose-adjusted warfarin in the Randomized Evaluation of Long-term Anticoagulation Therapy (RE-LY) trial^[86] in doses of 110 and 150 mg twice daily in approximately 12000 patients, 20% of whom had a history of TIA or stroke. The lower dose of dabigatran was non-inferior to warfarin in reducing the rate of stroke or systemic embolism, while the higher dose was superior. Both doses significantly reduced major bleedings compared with warfarin in patients younger than 75 years, whereas in elderly patients the lower dose was associated with a similar rate and the higher dose with an increased rate of major bleedings^[87]. The factor Xa inhibitor rivaroxaban (20 mg daily) was compared with warfarin in the Rivaroxaban-Once daily, oral, direct factor Xa inhibition Compared with vitamin K antagonism for prevention of stroke and Embolism Trial in AF (ROCKET-AF)^[88]. The study population in this trial was at high risk of stroke: 55% of patients had previous stroke or TIA and 90% had either a previous stroke or TIA, or three or more risk factors for stroke (mean CHADS₂ score 3.5). Rivaroxaban was non-inferior to warfarin for the prevention of stroke or systemic embolism and had a similar adverse effect profile. In the Apixaban for Reduction in Stroke and Other Thromboembolic Events in AF trial^[89] apixaban was better than warfarin in reducing the rate of embolic events and was associated with significantly less major bleedings, less intracranial haemorrhages and lower mortality. Finally, the Apixaban *vs* Aspirin to Prevent Stroke (AVERROES) trial investigated apixaban (5 mg twice daily) versus aspirin (81-324 mg daily) in patients with AF who were unsuitable or unwilling to receive VKAs^[90].

Table 3 Anticoagulation therapy after stroke or transient ischaemic attack in patients with atrial fibrillation

	Recommendation	Alternatives in patients unsuitable to anticoagulation
AHA/ASA ^[25]	Warfarin (INR 2.0-3.0)	Aspirin
ESO ^[24]	Warfarin (INR 2.0-3.0)	Aspirin plus dipyridamole
ACCP ^[52]	Dabigatran (150 mg twice daily)	Aspirin plus clopidogrel

AHA/ASA: American Heart Association/American Stroke Association; ESO: European Stroke Organisation; ACCP: American College of Chest Physicians; INR: International normalized ratio.

The rate of stroke or systemic embolism was significantly lower with apixaban than with aspirin, whereas the rates of major bleeding, intracranial haemorrhage, gastrointestinal bleeding, and myocardial infarction were similar in both groups. Another large phase III trial with a new anticoagulant, edoxaban, is ongoing^[91]. The proportion of patients with AF who were enrolled in all these trials and had a previous stroke or TIA varied between 14% in AVERROES and 55% in ROCKET-AF. In each trial, the absolute rate of stroke or systemic embolism was higher in patients with a previous stroke or TIA than in those without, but the relative effect of each new anticoagulant compared to warfarin in the prevention of stroke or systemic embolism was consistent among patients with or without previous cerebral ischaemic events. Similarly, the relative effect of each new agent on major bleeding did not differ^[92-95]. Indirect comparisons suggest that the relative effects of each of the new oral anticoagulants compared with warfarin were consistent for stroke and systemic embolism, haemorrhagic stroke, and mortality, but not for myocardial infarction and extracranial major bleeding^[96-102]. Similar results have been confirmed in a recent meta-analysis of data from 12 phase II and phase III randomized controlled trials comparing new oral anticoagulants with warfarin^[103].

The majority of current guidelines still regard VKAs at INR 2.0-3.0 to be the standard treatment after cerebral ischaemia of cardiac origin, including AF, for patients who are suitable for such therapy, with few exceptions, on the basis of recent findings concerning new anticoagulants^[24,25,52] (Table 3). Although these drugs have some advantages compared to warfarin, the unknown long-term safety profile, the absence of an antidote in the case of bleeding and their costs are important issues to consider.

Antihypertensive agents

Long term management of hypertension significantly reduces the risk of recurrent events. A meta-analysis of 7 randomized controlled trials including 15527 patients with TIA or stroke randomly allocated to treatment group within 1-14 mo after the event, showed reduction of recurrent stroke, myocardial infarction and vascular death^[38]. Benefit was present irrespective of a previous diagnosis of hypertension and risk reduction was greater with larger BP lowering. However, there is uncertainty regarding the class of drugs to prefer because of the small number of trials included. Significant reductions in recurrent stroke were seen with diuretics alone and in combination with angiotensin-converting enzyme inhibitors

(ACEIs) but not with β -blockers (BBs) or ACEIs used alone; nonetheless, statistical power was limited, particularly for the assessment of β -blockers, and calcium channel blockers (CCBs) and angiotensin receptor blockers (ARB) were not evaluated in any of the included trials. One of the largest trial included in the meta-analysis, the PROGRESS trial, showed a 26% reduction of the risk of stroke with the combination of perindopril and indapamide compared to placebo^[104]. Patients were included regardless of baseline BP and benefits were seen even in those patients who were normotensive before admission. Since this meta-analysis, 2 additional large-scale randomized trials were published: the Morbidity and Mortality after Stroke, Eprosartan Compared to Nitrendipine for Secondary Prevention (MOSES) trial^[105], and the PROFESS trial^[106]. In the MOSES, patients with an ischaemic event in the previous 2 years were randomly allocated to eprosartan (an ARB) or nitrendipine (a CCB). The risk of stroke was lower in the former group, but the benefit was only due to fewer TIAs, with no significant effect on stroke^[105]. The PROFESS failed to show a protective role of telmisartan, an ARB, in stroke prevention. Similarly, another study found no difference in a composite outcome of death from vascular causes, myocardial infarction, stroke or hospitalization for heart failure between patients with previous vascular event of any kind or diabetes randomly allocated to telmisartan or ramipril^[107]. The Telmisartan to Prevent Recurrent Stroke and Cardiovascular Events (TRANSCEND) trial^[108] showed only a modest effect on prevention of vascular events in patients treated with telmisartan, thus confirming the lack of evidence of ARB effectiveness in stroke prevention.

On the basis of the available evidence, current guidelines recommend antihypertensive treatment in most patients with recent TIA or stroke, beyond the first 24 h, irrespective of a history of hypertension before admission, with some favouring the combination of an ACEI and a diuretic agent, on the basis of PROGRESS trial^[25]. An absolute BP target and reduction are uncertain and should be individualized, but benefit has been associated with an average reduction of roughly 10/5 mmHg, and normal BP levels have been defined as $> 120/80$ mmHg^[109].

However, recent studies have shown that visit-to-visit variability in BP and episodic hypertension are powerful risk factors for stroke, independently of mean BP, and that the benefits of some BP-lowering drugs are attributable partly to reduce variability in BP levels^[110,111]. Although reductions in mean SBP are similar with different drug classes, CCBs and diuretics also reduce variability

in SBP, while BBs increase variability^[112,113]. In particular, it has been shown that higher doses of CCBs reduce BP variability more than lower doses, and higher doses of BBs increase BP variability^[114]. This means that combinations containing a low dose of BBs will have less adverse effects on variability in SBP and the associated risk of stroke, but low doses of CCBs will be less effective in reducing BP variability. Drug class effects on variability persist even in combination therapy.

Another issue to consider is the heterogeneity of stroke patients. Elderly patients seem to benefit most from a combination therapy of ACEIs and diuretics^[115], as well as patients with renal impairment or diabetic nephropathy, in whom ACEIs and ARBs are recommended. The detrimental effect of such drugs on BP variability could be offset by the combination use with CCBs, as suggested in the Avoiding Cardiovascular Events through Combination Therapy in Patients Living with Systolic Hypertension study^[116], in whom patients with hypertension and previous vascular disease, including stroke, were randomly allocated to a combination of benazepril plus amlodipine or a combination of benazepril plus hydrochlorothiazide. The former group of patients had fewer combined cardiovascular deaths and chronic kidney disease events compared to the latter.

However, direct comparisons of drugs for secondary prevention of stroke are needed in order to recommend the most appropriate BP-lowering drug regimen. More research is required to identify ideal combinations and doses of drugs for the prevention of stroke in specific patient groups with particular reference to their effect on BP variability.

Lipid lowering agents

Although the association between serum lipid concentrations and ischaemic risk is weaker for stroke than for myocardial infarction^[117], lipid levels modification is important in both primary and secondary prevention of stroke^[118]. A recent observational study^[144] showed a better functional outcome in stroke patients who received statin therapy before admission to hospital and who continued such a treatment during hospitalization. The prognosis was even better with higher doses of statins and when therapy was started earlier. Regarding secondary prevention, in a retrospective analysis of the Heart Protection Study (HPS)^[119] restricted to subjects with a history of stroke, simvastatin did not significantly reduce the risk of stroke. In contrast, the Stroke Prevention by Aggressive Reduction in Cholesterol Levels (SPARCL) trial^[120] showed a 16% reduction in stroke recurrence in patients with previous stroke or TIA, not of cardiac origin, randomized to atorvastatin 80 mg daily versus placebo, within 6 mo from index event. One concern in the SPARCL trial and in the subset of patients in HPS^[119] who had a stroke before randomization was a significant increase in the risk of haemorrhagic stroke in patients allocated to statin treatment^[121]. Further analysis showed that much of excess risk of haemorrhagic stroke on atorvastatin therapy was in patients with a small vessel disease stroke

as qualifying event, either of haemorrhagic or ischaemic type^[122]. Although in the former group there was no overall benefit, in the latter the net benefit was still in favour of treatment because of a reduction in ischaemic recurrences^[122]. This suggests that statin therapy should be initiated only in patients with atherosclerotic stroke or in presence of other noncerebrovascular indications, such as coronary heart disease or diabetes^[123].

Another important clinical issue relates to low-density lipoprotein (LDL) concentration. In a meta-analysis of all statin trials that looked at intense LDL lowering versus standard treatment, achievement of an LDL concentration of 100 mg/dL resulted in 16% relative reduction in risk of stroke. Subanalysis of the SPARCL trial showed a 28% reduction in stroke risk with a LDL concentration lower than 70 mg/dL in comparison with a level of 100 mg/dL^[124,125]. The Treat Stroke to Target trial^[126] is an ongoing study aimed at unravelling the issue regarding the optimal LDL target to reach. The working hypothesis of this trial is that the achievement of an LDL concentration of 70 mg/dL is better than 100 mg/dL in patients with previous atherosclerotic TIA or stroke. The Optimal BP and Cholesterol Targets for Preventing Recurrent Stroke in Hypertensive (ESH-CHL-SHOT) trial^[127] will be started soon to address this issue. Its purpose, indeed, is to investigate both SBP and LDL cholesterol levels to achieve to optimize secondary prevention. Regarding lipid-lowering treatment in particular, the patients will be randomly allocated to one of two different LDL targets, 70 to 110 mg/dL or < 70 mg/dL, using different statins, including simvastatin, pravastatin, fluvastatin, atorvastatin or rosuvastatin.

Furthermore, there is still uncertainty regarding whether to start with high-dose or low-dose statin therapy, considering benefit and potential side effects. The SPARCL trial^[120] reliably showed that high-dose atorvastatin at 80 mg/d reduces the risk of stroke and other cardiovascular recurrences in patients with non cardioembolic stroke. It is however unclear, in the absence of randomized comparisons in patients with stroke, if lower doses of atorvastatin are equally or less effective than the high-dose regimen. There is evidence that atorvastatin at 80 mg/d provides significant reductions in the risk of cardiovascular events and stroke beyond that provided by a 10-mg dose in patients with coronary disease^[128]. However, the high-dose regimen was complicated more frequently by higher rates of adverse events and discontinuation, both in Treating to New Targets (TNT) trial^[128] and SPARCL trial^[120]. Moreover, the Food and Drug Administration issued a warning regarding increased risk of myopathy in patients taking simvastatin at 80 mg/d, especially if taken in combination with CCBs, which are frequently used in patients with stroke. Although the SPARCL results support the use of high-dose atorvastatin for secondary prevention of stroke, it is unclear whether the highest currently approved doses of statins have neuroprotective effects. It is also unknown if other hydroxymethylglutaryl-coenzymeA reductase inhibitors, in different doses, could be potentially effective in pre-

vention of stroke recurrences. In this regard, two trials are ongoing to investigate the combined effect of this class of drugs^[127] and that of pravastatin^[129] in particular in long-term secondary prevention after stroke.

On the basis of available data the American Heart Association/American Stroke Association (AHA/ASA) guidelines recommend statin therapy with intensive lipid-lowering effect for secondary prevention among patients with ischemic stroke or TIA who have evidence of atherosclerosis, an LDL-level ≥ 100 mg/dL, and who are without known coronary heart disease. They suggest, as a target, a reduction of at least 50% in LDL or, alternatively, an LDL level of < 70 mg/dL^[25].

Other medications used to treat dyslipidemia include niacin, fibrates, and cholesterol absorption inhibitors. These agents can be used by patients intolerant to statins, but the evidence of their efficacy for prevention of stroke recurrence is scarce. Niacin has been associated with a reduction in cerebrovascular events^[130], whereas gemfibrozil reduced the rate of total strokes among men with coronary artery disease and low levels of high-density lipoprotein cholesterol (≤ 40 mg/dL) in the Veterans Affairs HDL Intervention Trial. The latter result was no more significant when adjudicated events alone were analyzed^[131].

Surgical procedures

Percutaneous PFO closure: PFO is a common embryonic defect of the interatrial septum, being present in up to 25% of healthy people. In a meta-analysis PFO was found to be significantly associated with increased risk of stroke in younger patients with cryptogenic stroke, especially when atrial septal aneurysm (ASA) is also present^[132]. This leads to the assumption that PFO may be the cause somehow associated to stroke occurrence, although it may also be an incidental finding^[133]. The role of PFO in determining the risk of recurrence is uncertain, as well. In a substudy of the PFO in Cryptogenic Stroke Study (PICSS)^[134] there were no differences in the rates of recurrent strokes in those with or without PFO, as well as no demonstrated effect on outcomes based on PFO size or presence of ASA. In contrast, the study of Mas *et al.*^[135] reported higher recurrence rates of stroke associated with both PFO and ASA, indicating that the presence of both these cardiac abnormalities was a predictor of an increased risk of recurrent stroke, whereas isolated PFO, whether small or large, was not. Thus, the natural history after cerebrovascular events in patients with PFO remains insufficiently defined. As a consequence, the optimal management strategy for treating patients with cryptogenic stroke who are discovered to have a PFO remains to be determined. Observational studies on medical treatment in patients with PFO with either antiplatelets or warfarin reported a risk of recurrent stroke or TIA ranging from 3% to 12% during the first year^[136]. Both larger PFO size and a greater degree of right-to-left shunt increased the risk of paradoxical embolism in one study^[136]. Of note, observational data for the comparative effectiveness of anticoagulants versus antiplatelet agents

for secondary prevention after stroke associated to PFO show a significant benefit of warfarin, whereas the only relevant randomized study, coming from a subgroup of PICSS^[134], found a non-significant trend in favor of warfarin over aspirin. On the basis of this evidence, AHA guidelines state that there are insufficient data to establish the superiority of warfarin over aspirin for secondary stroke prevention after stroke or TIA in patients with PFO^[25]. However, warfarin might be more beneficial in presence of deep venous thrombosis or ASA, according to the European Stroke Organization guidelines^[24]. Another therapeutic possibility is percutaneous PFO closure, which has supplanted surgical procedures. Case series and nonrandomized comparisons have long suggested that closure is a highly efficacious procedure and have led to the rapid off-label adoption of this intervention^[137]. In contrast, recent randomized controlled trials failed to identify any statistically significant difference between closure and medical treatment for secondary prevention after stroke^[138-140]. However, such trials have several limitations and low statistical power. In the Evaluation of the STARFlex Septal Closure System in Patients with a Stroke and/or TIA due to Presumed Paradoxical Embolism through a PFO (CLOSURE) trial^[138] the main limitation was the low outcome rate, compared to previous observational studies, thus pointing to a failure to select patients with PFO-related events. This may, in part, be attributable to a reluctance to randomize those patients with cryptogenic stroke who appear most likely to have had a PFO-related event (*i.e.*, patients with clinical indicators of paradoxical embolism, such as large shunt, associated ASA, Valsalva maneuver at onset of stroke, or absence of any conventional stroke risk factors). Other limitations of this trial include idiosyncratic effects, such as *in situ* thrombosis or other mechanical complications, with the use of the obsolete STARFlex device, which may have increased the outcome rates in the intervention arm, as well as the short follow-up, that may be inadequate to capture the benefit of the intervention^[137]. The Randomized Evaluation of Recurrent Stroke Comparing PFO Closure to Established Current Standard of Care Treatment (RESPECT) trial^[139] confirmed the results of the CLOSURE trial in intention-to-treat analysis, in spite of important differences between the two studies with respect to the study design, the population included, and the device tested. In the RESPECT trial^[139] the enrollment criteria were more stringent and the follow-up period was longer than in the CLOSURE. Moreover, the Amplatzer PFO Occluder, as compared to STARFlex used in the CLOSURE, was associated with higher effective closure rate. Finally, although different in terms of population and endpoints, the Percutaneous Closure of PFO in Cryptogenic Stroke (PC) trial^[140] corroborated previous results. However, unlike the results of the CLOSURE trial, the results of the RESPECT and the PC trials have encouraged the supporters of PFO closure. The advocates of closure focus on the modest statistical power of these trials, the substantial relative effect size of the point estimates in both trials, the signifi-

cance of the per-protocol and as-treated analyses in the RESPECT, and arbitrariness of the conventional *P*-value threshold of 0.05. Although the controversy over efficacy may not be settled, it is worth noting that the safety profile of Amplatzer device appeared to be superior to that of STARFlex device tested in the CLOSURE. This supports the hypothesis that PFO closure, in certain conditions, could be superior to medical treatment and reinforces the need to carry on trials with more selected populations. Indeed, other trials are ongoing, such as the PFO Closure or Anticoagulants Versus Antiplatelet Therapy to Prevent Stroke Recurrence (CLOSE) trial^[141] and the GORE® HELEX® Septal Occluder/GORE® Septal Occluder for PFO Closure in Stroke Patients (REDUCE) trial^[142]. However, like previous analyses, these new studies are hampered by a slow and skewed enrollment since most patients assumed to be at high risk undergo percutaneous closure. Besides, stroke pathophysiology is multifactorial, and the diagnosis of PFO-mediated paradoxical embolism is presumptive. Both a PFO and a cryptogenic stroke may coexist without causal relation in a given patient. In this case, obviously, PFO closure will not reduce the risk of recurrence. Current guidelines, which are antecedent to these recent trials, do not recommend closure because of the paucity of data^[25]. However, pending other trials and based on RESPECT, it would be reasonable to discuss closure with Amplatzer device in young patients (< 50 years) with a substantial shunt and a cortical infarct, in the absence of known vascular risk factors^[143].

Carotid endarterectomy and stenting for large-artery atherosclerosis

Beside medical therapy, the mainstays of treatment options for symptomatic large-artery atherosclerosis are carotid endarterectomy (CEA) and carotid angioplasty and stenting (CAS), the latter being emerged more recently as an alternative to surgical intervention to prevent recurrent strokes.

CEA plus medical therapy is more effective and safe than medical therapy alone in secondary prevention after stroke for symptomatic patients with > 70% carotid stenosis according to 3 major trials, the European Carotid Surgery Trial^[144], the North American Symptomatic CEA Trial^[145], and the Veterans Affairs Cooperative Study^[146]. However, in all these trials medical therapy did not include aggressive atherosclerotic medical management based on statins, antihypertensive agents, antiplatelets other than aspirin, and smoking cessation, thus meaning that the benefit of surgery today may be less than in the early trials. Uncertainty exists for patients with symptomatic stenosis in the range of 50% to 69%, for whom CEA should be considered only with appropriate case selection when the risk-benefit ratio is favorable for the patient. The timing of CEA after an acute event is controversial, with experts advocating waiting anywhere from 2 to 6 wk^[25]. However, early surgery (< 2 wk) turned out to be beneficial, according to pooled analysis of trials. Furthermore, early intervention (< 3 wk) may be beneficial and

safe in low-risk patients with TIA and minor strokes, and in those without haemorrhagic transformation^[25].

CAS is a percutaneous, less invasive procedure that has emerged as an interesting alternative to CEA for the treatment of extracranial carotid artery disease, especially thanks to the advances in endovascular technology, including embolic protection devices (EPD) and improved stent design. Complication rates are comparable to CEA. Although advantages related to the less invasive procedure, CAS durability remains unproven. The largest randomized trial in symptomatic patients, the Carotid and Vertebral Artery Transluminal Angioplasty Study 2 trial^[147], showed comparable outcome between the two procedures, also in the long-term follow-up. The Stent-Supported Percutaneous Angioplasty of the Carotid Artery Versus Endarterectomy trial^[148] which was ended after the second interim analysis owing to the low rate of recruitment, showed similar rates of periprocedural stroke or death and ipsilateral ischaemic stroke up to 2 years between CEA and CAS. However, the fact that only 27% of patients with CAS in this study used EPD may have skewed the outcome. Because EPD reduce periprocedural stroke rates, they have been required in all stenting trials since then. The Stenting and Angioplasty with Protection in Patients at High Risk for Endarterectomy trial^[149] randomized both symptomatic and asymptomatic high risk patients, showing that CAS was not inferior to CEA. However, a high percentage of patients were asymptomatic (70%), thus limiting the applicability of the results to symptomatic ones. The Endarterectomy Versus Angioplasty in Patients with Symptomatic Severe Carotid Stenosis trial^[150] showed higher rates of periprocedural stroke or death at 30 d and within 4 years, as well. However, the results of this trial are influenced by the limited experience of the operators, the use of five different stents and seven different cerebral protection devices, and the fact that EPD were optional in the early phase of the study. The most important trial on this topic is the Carotid Revascularization Endarterectomy Versus Stenting Trial^[151], which included patients with both symptomatic and asymptomatic carotid disease in similar proportions. It demonstrated that both procedures had similar safety and efficacy profiles for all patients, according to gender and the presence or not of previous cerebrovascular events. However, patients aged 70 years or older had better outcomes with CEA for the primary endpoint (any stroke, death or myocardial infarction (MI) within 30 d and ipsilateral stroke during long-term follow-up) and less adverse events. Younger patients had greater benefit from CAS than older subjects. The CAS group had a greater percentage of patients with stroke within the first 30 d after the procedure, but a lower rate of MI events. In trials of symptomatic patients, stenting was associated with a significantly higher risk of 30-d stroke or death, particularly in people over 70 years, compared to CEA, whereas the long-term rate of ipsilateral stroke were similar for both interventions^[152]. However, stenting was associated with less MI and cranial nerve palsies.

On the basis of these studies, current AHA/ASA

guidelines recommend that patients who have experienced a TIA or stroke within the past 6 mo and have ipsilateral carotid stenosis of 70%-90% would be candidates for CEA if the perioperative morbidity and mortality risk is less than 6%. In symptomatic patients with a 50%-60% stenosis, the decision to perform CEA depends on factors such as age, comorbidities, and sex. If a patient is a good candidate for CEA, intervention within 2 wk is recommended^[25]. To date, CAS is considered an alternative option for symptomatic patients with carotid stenosis of more than 70% as determined by noninvasive imaging or more than 50% as determined by catheter angiography. CAS is also an alternative in patients deemed at high risk for surgical intervention. High risk is defined by presence of severe comorbidities and challenging anatomical features (such as prior neck intervention or irradiation, postendarterectomy restenosis, surgical inaccessible lesion, contralateral carotid occlusion, contralateral vocal cord palsy, or the presence of tracheostomy)^[25]. Patients are also recommended to optimize medical therapy with antiplatelets, statins and risk factor modifications.

Angioplasty and stenting for intracranial atherosclerosis

According to the Stenting versus Aggressive Medical Therapy for Intracranial Arterial Stenosis (SAMMPRIS) trial^[153], angioplasty and stenting of intracranial stenosis did not reduce the risk of recurrent stroke compared with aggressive medical therapy, including dual antiplatelets, statins and antihypertensive agents.

CONCLUSION

Secondary prevention with antiplatelet and antihypertensives medications, statins and anticoagulants as appropriate, should be initiated urgently after TIA or minor strokes because of the high risk of early stroke recurrence. For long-term prevention, aspirin plus dipyridamole or clopidogrel must be proposed as first-line approach after cerebral ischaemia of arterial origin. For cardioembolic strokes in patients with non-valvular AF new anticoagulation agents are challenging the current standards of VKAs. Lipid and blood-pressure-lowering treatments are warranted in all cerebral ischaemia, but recommendations about optimal drug regimen, including doses and when to start therapy, as well as target levels of LDL or BP to be achieved, are still debated. Revascularization procedures, including CEA or stenting, must be taken into consideration early after the event in patients with stroke or TIA related to carotid atherosclerotic disease.

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Molecular diagnosis of autosomal recessive cerebellar ataxia in the whole exome/genome sequencing era

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Abstract

Autosomal recessive cerebellar ataxias (ARCA) are a clinically and genetically heterogeneous group of rare neurodegenerative disorders characterized by autosomal recessive inheritance and an early age of onset. Progressive ataxia is usually the prominent symptom and is often associated with other neurological or additional features. ARCA classification still remains controversial even though different approaches have been proposed over the years. Furthermore, ARCA molecular diagnosis has been a challenge due to phenotypic overlap and increased genetic heterogeneity observed within this group of disorders. Friedreich's ataxia and ataxia telangiectasia have been reported as the most frequent and well-studied forms of ARCA. Significant progress in understanding the genetic etiologies of the ARCA has been achieved during the last 15 years. The methodological revolution that has been observed in genetics over the last few years has contributed significantly to the molecular diagnosis of rare diseases including the ARCAs. Development of high throughput technologies has resulted in the identification of new ARCA genes and novel mutations in known ARCA genes. Therefore,

an improvement in the molecular diagnosis of ARCA is expected. Moreover, based on the fact that many patients still remain undiagnosed, additional forms of ataxia are expected to be identified. We hereby review the current knowledge on the ARCAs, focused on the genetic findings of the most common forms that were molecularly characterized before the whole exome/genome era, as well as the most recently described forms that have been elucidated with the use of these novel technologies. The significant contribution of whole-exome sequencing or whole-genome sequencing in the molecular diagnosis of ARCAs is discussed.

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Key words: Autosomal recessive cerebellar ataxia; Whole-exome sequencing; Whole-genome sequencing; Homozygosity mapping; Next generation sequencing

Core tip: Molecular diagnosis of autosomal recessive cerebellar ataxias (ARCA) is challenging due to clinical overlap and increased genetic heterogeneity. Although use of traditional techniques led to the identification of causative mutations in the past, the recent employment of novel technologies in this field, has initiated a new era in the molecular diagnosis of ARCA. Limitations such as small sized families, large numbers of candidate genes within mapped intervals and large sized genes hindered the timely discovery of ARCA genes using conventional Sanger sequencing. ARCA gene discovery and molecular diagnosis should be achievable at a much faster rate through the use of whole-genome or whole-exome sequencing technologies.

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INTRODUCTION

Autosomal recessive cerebellar ataxias (ARCA) constitute a subgroup of the hereditary cerebellar ataxias^[1]. They are clinically and genetically heterogeneous neurodegenerative disorders characterized by autosomal recessive inheritance and usually a clinical onset before the age of 20. Both the central and the peripheral nervous systems may be involved, as well as other systems and organs in rare cases^[1,2]. In most cases the cerebellum, the spinocerebellar tract and/or the sensory tracts of the spinal cord are primarily affected^[3]. Therefore, progressive ataxia is the main symptom and is often associated with other neurological or extra-neurological signs.

This group encompasses many rare diseases; Friedreich's ataxia and ataxia telangiectasia being the most frequent forms^[2-4]. Their prevalence has been estimated to be 7 in 100000 inhabitants^[5] with considerable variability observed in different geographic regions^[3,6,7]. Many classifications have been proposed using different criteria and this issue still remains controversial. The first clinical classification of inherited ataxias, also encompassing ARCA, was proposed by Harding in 1983 and was based on the age of onset and pathological mechanisms^[8,9]. Through the next decade, genetic studies led to new insights and thus additional criteria were used in some classifications. Koenig^[10] proposed a classification of ARCA based on topographical and pathophysiological criteria, while in the next year the group of Filla proposed a pathogenic classification of the hereditary ataxias^[11]. More recently, Palau and Espinos proposed a classification of ARCA that is based on inheritance patterns and natural history of the clinical symptoms^[1].

Through the past 15 years, significant progress has been made in improving our understanding of the genetic etiology of the ARCA. Currently, more than 30 genes have been associated with ARCA (Table 1) and many more are expected to be discovered since the genetic causes still remain unknown, for a large number of patients/families. The identified ARCA genes are involved in variable cellular processes such as mitochondrial energy generation, DNA repair, transcription, RNA processing, protein folding and ion channels^[12].

Despite the existence of different classification schemes and the significant number of already identified genes, diagnosis of ARCA remains a challenge due to their vast clinical variability and genetic heterogeneity^[13-15]. The methodological revolution of genetics observed over the last years contributed significantly to the proper diagnosis of ARCA. The recently developed next-generation sequencing (NGS) technologies, whole-exome sequencing (WES)/whole-genome sequencing (WGS), or the strategy of using homozygosity linkage mapping combined with NGS, proved to be efficient for the identification of novel genes associated with rare diseases. In comparison to the traditional methods widely used in the past for gene identification, the new techniques are easier, cost and time-efficient since they offer the opportunity to screen the whole-genome or the coding part of the genome in

one experiment^[14]. Limitations such as locus heterogeneity, small family size and existence of numerous candidate genes within a mapped region are no longer an obstacle^[16]. Although relatively new, these technologies proved to be powerful methods for gene identification in rare neurological disorders such as ataxia; 7 new genes were identified in the past 3 years. Moreover, exome sequencing is gradually used as a screening tool thus enabling the identification of mutations in already known genes^[17-21].

We hereby provide a review focused on the molecular diagnosis of ARCA with more emphasis on the recent discoveries observed in the whole exome/genome sequencing era.

MOLECULAR DIAGNOSIS OF AUTOSOMAL RECESSIVE CEREBELLAR ATAXIAS WITH THE CONVENTIONAL METHODS

The most common ARCA forms that have been molecularly diagnosed through the use of traditional methods as well as the most common mutations identified are outlined below. If an affected individual is clinically suspected of having a specific type of the following ARCA, then investigation could begin with targeted mutation analysis.

Degenerative ataxias

Friedreich ataxia: Friedreich ataxia is the most frequent inherited ataxia in the Caucasian population^[1] and the most common form of degenerative recessive ataxia with an estimated prevalence of 1-2/50000^[22-24]. The disease has been associated with mutations in the *FXN* gene located on chromosome 9q13^[25-27]. A homozygous GAA repeat expansion within the first intron of the gene is the causative mutation for 95% of the cases^[28], while the rest 5% of the cases are compound heterozygous for a GAA expansion and a point mutation in the *FXN* gene^[29]. The expansion mutation leads to an impairment of transcription and further to a significant reduction in the expression of the encoded protein, frataxin. Although the function of frataxin has not been fully elucidated, its role in cellular iron homeostasis has been established. The deficiency caused by the expansion mutation results in mitochondrial dysfunction and oxidative damage^[24,30].

Friedreich ataxia is considered as the first candidate for genetic testing of patients with ARCA in most populations. Specific polymerase chain reaction (PCR)-based molecular tests for the GAA repeat expansion are available in many laboratories^[31-33]. Suspected compound heterozygote cases may be further investigated by Sanger sequencing of the *FXN* gene coding regions.

Mitochondrial recessive ataxia syndrome: Mitochondrial recessive ataxia syndrome is the most frequent juvenile- or adult-onset type of ataxia in Finland^[34]. The disease has been associated with mutations in the *POLG* gene, encoding polymerase γ which is implicated in

Table 1 Autosomal recessive cerebellar ataxias

Type	Gene/protein	Gene OMIM#	Gene locus
Degenerative ataxias			
Friedreich's ataxia	<i>FXN</i> /Frataxin	*606829	9q13
SCAR9	<i>ADCK3</i> /aarF domain-containing protein kinase 3	*606980	1q42.13
MIRAS	<i>POLG</i> /Polymerase γ	*174763	15q25
IOSCA (MTDPS7)	<i>C10orf2</i> /Twinkle	*606075	10q24
Marinesco-Sjögren syndrome	<i>SIL1</i> /BiP associated protein	*608005	5q31
Charlevoix-Saguenay spastic ataxia	<i>SACS</i> /Sacsin	*604490	13q12
EOCARR (EOCA)	-	-	13q11-12
Ataxias with DNA repair defects			
AOA1	<i>APTX</i> /Aprataxin	*606350	9p13.3
AOA2 (SCAR1)	<i>SETX</i> /Senataxin	*608465	9q34
AOA3	<i>PIK3R5</i> /Phosphoinositide 3-kinase regulatory subunit 5	*611317	17p13.1
AT	<i>ATM</i> /ATM	*607585	11q22.3
ATLD	<i>MRE11A</i> /MRE11	*600814	11q21
SCAN1	<i>TDP1</i> /Tyrosyl DNA phosphodiesterase 1	*607198	14q31-q32
Congenital ataxias			
JBTS1	<i>INPP5E</i> /Phosphatidylinositol polyphosphate 5-phosphatase type IV	*613037	9q34.3
JBTS2	<i>TMEM216</i> /transmembrane protein 216	*613277	11p12-q13.3
JBTS3	<i>AHI1</i> /Joubertin	*608894	6q23.3
JBTS4	<i>NPHP1</i> /Nephrocystin-1	*607100	2q13
JBTS5	<i>CEP290</i> /Nephrocystin-6	*610142	12q21.3
JBTS6	<i>TMEM67</i> /Meckelin	*609884	8q22.1
JBTS7	<i>RPGRIPL1</i> /Nephrocystin-8	*610937	16q12.2
Cayman ataxia	<i>ATCAY</i> /Caytaxin	*608179	19p13.3
Metabolic ataxias			
AVED (VED)	α -TTP/ α -tocopherol transfer protein	*600415	8q13.1-q13.3
ABL	<i>MTP</i> /Microsomal triglyceride transfer protein	*157147	4q22-24
CTX	<i>CYP27</i> /Sterol 27-hydroxylase	*606530	2q33-qter
Refsum's disease	<i>PHYH</i> /Phytanoyl-CoA hydrolase	*602026	10pter-p11.2
	<i>PEX7</i> /Peroxin 7	*601757	6q22-q24
Metachromatic leucodystrophy	<i>ARSA</i> /Arylsulfatase 1	*607574	22q13.31-qter
Niemann Pick disease type C	<i>NPC1</i> /NPC1 protein	*607623	18q11-q12
GM1 gangliosidosis	<i>GLB1</i> /Beta-galactosidase	*611458	3p21.33
GM2 gangliosidosis	<i>HEXA</i> /Hexosaminidase A	*606869	15q23-24
Wilson disease	<i>ATP7B</i> /ATPase Cu transporting beta-polypeptide	*606882	13q14.3
Aceruloplasminemia	<i>CP</i> /Ceruloplasmin	*117700	3q23-q24
CHAC	<i>VPS13A</i> /Chorein	*605978	9q21
Other recently identified types			
SCAR8	<i>SYNE1</i> /Syne-1	*608441	6q25.1-q25.2
Rundataxin-related ataxia	<i>KIAA0226</i> /Rundataxin	*613516	3q27.3-qter
Novel types identified by the use of next-generation sequencing			
SCAR10	<i>ANO10</i> /Anoctamin 10	*613726	3p22.1
SCAR11	<i>SYT14</i> /Synaptotagmin XIV	*610949	1q32.2
SCAR7	<i>TPP1</i> /tripeptidyl-peptidase 1 enzyme	*607998	11p15.4
Autosomal-recessive cerebellar ataxia with spasticity	<i>GBA2</i> / β -glucosidase 2	*609471	9p13.3
SCAR13	<i>GRM1</i> /Metabotropic glutamate receptor 1	*604473	6q24.3

SCAR7: Spinocerebellar ataxia autosomal recessive type 7; SCAR8: Spinocerebellar ataxia autosomal recessive type 8; SCAR9: Spinocerebellar ataxia autosomal recessive type 9; SCAR10: Spinocerebellar ataxia autosomal recessive type 10; SCAR11: Spinocerebellar ataxia autosomal recessive type 11; SCAR13: Spinocerebellar ataxia autosomal recessive type 13; MIRAS: Mitochondrial recessive ataxic syndrome; IOSCA: Infantile-onset spinocerebellar ataxia; AOA: Ataxias with oculomotor apraxia; AT: Ataxia telangiectasia; ATLD: Ataxia-telangiectasia-like disorder; SCAN1: Spinocerebellar ataxia with axonal neuropathy; CTX: Cerebrotendinous xanthomatosis; CHAC: Chorea-acanthocytosis; JBTS: Joubert syndrome; AVED: Ataxia with isolated vitamin E deficiency; ABL: Abetalipoproteinemia.

mtDNA replication and repair. The most common mutation identified thus far is the c.2243G > C [p.Trp748Ser] associated with the c.3428A > G [p.Glu1143Gly] mutation which is probably a polymorphism with a modifying role. Another frequent mutation is the c.1399G > A [p.Ala467Thr]. Homozygous or compound heterozygous patients for the three mutations have been reported^[34-36]. The presence of the three mutations may be molecularly diagnosed by PCR and solid-phase minisequencing while

a Real time PCR assay has been proposed for identifying the presence of the c.1399G > A mutation^[34].

Infantile onset spinocerebellar ataxia: Infantile onset spinocerebellar ataxia is a rare disorder manifesting at a very early age and was first described in Finland. It has been associated with mutations in the *C10orf2* gene on chromosome 10q24 that encodes Twinkle, a mtDNA specific helicase and Twinky, a rarer splice variant^[37]. The most com-

mon mutation identified is c.1523A > G [p.Tyr508Cys]. Mutations in this gene may be detected either by direct sequencing or by solid-phase minisequencing.

Autosomal recessive spastic ataxia of Charlevoix-Saguenay: Autosomal recessive spastic ataxia of Charlevoix-Saguenay (ARSACS) is an early childhood onset disease characterized by spastic gait ataxia. It was first described in families from the Charlevoix-Saguenay region of northeastern Quebec and for many years it was thought to be restricted there. It has been mapped to chromosome 13q11 and causative mutations have been identified in the *SACS* gene^[38] that encodes saccin, a chaperone protein. Almost 96% of the molecularly diagnosed individuals from this region are homozygote or compound heterozygote for the two founder mutations c.8844delT [p.Pro2948fs] and c.7504C > T [p.Arg2502*] (originally reported as 6594delT and 5254C > T) identified by sequence-based analyses^[39,40]. More than 100 other pathogenic mutations in individuals outside Quebec were identified since mutation analysis became available, demonstrating that mutations in *SACS* are a frequent cause of the disease worldwide^[41-43]. Molecular genetic testing is usually performed by targeted mutation analysis, by direct sequencing of all coding exons and their flanking intronic sequences and by deletion/duplication analysis using the multiplex ligation-dependent probe amplification method^[44].

Marinesco-Sjögren syndrome: Marinesco-Sjögren syndrome is a rare disorder with onset in infancy or childhood. Homozygosity mapping in two large consanguineous families of Turkish and Norwegian origin mapped the disease gene on chromosome 5q31^[45]. Causative mutations have been identified in patients of different ethnic origin, in the *SIL1* gene encoding a protein that acts as a nucleotide exchange factor for the heat-shock protein 70 chaperone the 78-kDa glucose-regulated protein, mainly located at the endoplasmic reticulum^[46-48]. Molecular genetic testing is performed by direct sequencing of all coding exons and their flanking intronic sequences.

Spinocerebellar ataxia autosomal recessive type 9: Spinocerebellar ataxia autosomal recessive type 9 (SCAR9), also known as autosomal recessive cerebellar ataxia type 2 (ARCA2) refers to ARCA associated with muscle coenzyme Q₁₀ deficiency^[49]. Homozygosity mapping (GeneChip Human Mapping 10K 2.0 Xba Array, Affymetrix) in a large consanguineous Algerian family mapped the disease gene to chromosome 1q41-q42. *ADCK3* gene mutations have been identified by direct sequencing in this family, in additional families and also in sporadic patients^[49-51]. The *ADCK3* protein is a mitochondrial ancestral kinase which is involved in CoQ10 synthesis^[49].

Ataxias with DNA repair defects

ARCA with oculomotor apraxia: ARCA with oculomotor apraxia (OMA), known as ataxias with oculomo-

tor apraxia (AOA), are characterized by cerebellar ataxia accompanied by ophthalmological and neurological symptoms^[52,53]. Five distinct genetic forms have thus far been identified: ataxia-telangiectasia (A-T), ataxia-telangiectasia-like disorder (ATLD), ataxia with oculomotor apraxia type 1 (AOA1), ataxia with oculomotor apraxia type 2 (AOA2) and ataxia with oculomotor apraxia type 3 (AOA3)^[54].

(1) Ataxia telangiectasia: Ataxia telangiectasia is the second most common ARCA after Friedreich ataxia. It presents in early childhood with progressive cerebellar ataxia, while later ataxia may be accompanied by a multitude of symptoms such as choreoathetosis, dystonia, telangiectasia that is the guiding diagnostic feature, immunodeficiency, cancer predisposition, sinopulmonary infection, immunosensitivity and radiosensitivity^[1,14]. The disease gene *ATM* is located on chromosome 11q23^[55,56]. It is one of the largest genes, comprising 66 exons and encodes a member of the phosphoinositol-3 kinase family of proteins that are involved in cell cycle checkpoint control and the DNA repair through phosphorylation of substrates^[1]. More than 500 mutations have been identified, that are positioned throughout all the coding exons of the gene^[14]. Due to the large size of the gene, molecular diagnosis with the traditional methods has been difficult. A multiplex scanning method, detection of virtually all mutations-SSCP has been proposed however with reduced sensitivity^[57]. The newly developed NGS technologies are promising for the diagnosis of diseases such as A-T with associated mutations in large genes.

(2) Ataxia-telangiectasia-like disorder: Ataxia-telangiectasia-like disorder (ATLD) is a very rare disorder that resembles A-T in its clinical phenotype^[58] with absence of telangiectases and immune deficiency. It has been mapped to chromosome 11q21 and a small number of causative mutations have been identified in the *MRE11A* gene^[59-61]. The encoded protein interacts with RAD50 and NBS1 thus forming the core of the MRN (*MRE11-RAD50-NBS1*) complex which is involved in DNA repair pathways, in DNA recombination and in multiple cell-cycle checkpoints control^[60].

(3) Ataxia with oculomotor apraxia type 1: Ataxia with oculomotor apraxia type 1 (AOA1) was first described in Portuguese and Japanese families. The disease gene has been mapped to chromosome 9p13.3^[62] and mutations in the *APTX* gene have been identified^[63]. The encoded protein aprataxin, localizes to the nucleus^[64] and the mitochondria^[65] and it is involved in the nuclear and the mtDNA repair machinery. AOA1 is the most frequent type of ARCA in Japan and the second most frequent in Portugal after Friedreich ataxia. The insertion c.689insT and the missense c.617C > T [p.Pro206Leu] mutations, are more common in the Japanese and the nonsense c.837G > A [p.W279X] mutation is more common in Portuguese patients. More than 20 mutations including missense/nonsense or splice site mutations, small dele-

tions/insertions and gross deletions have been reported thus far in patients of different ethnicities, thus expanding the presence of AOA1 in other countries outside Japan and Portugal^[3,66-70].

(4) Ataxia with oculomotor apraxia type 2: Ataxia with oculomotor apraxia type 2 (AOA2) is also referred to as autosomal recessive spinocerebellar ataxia 1 (SCAR1) since oculomotor apraxia is an occasional finding, not present in all the patients. It is the second most frequent recessive ataxia in Europe^[4] and is usually (in 96% of cases) accompanied by a characteristic laboratory finding, the elevation of serum α -fetoprotein. AOA2 usually presents within the second decade of life and has a higher mean age of onset compared to AOA1 and A-T^[52]. The disease gene has been mapped to chromosome 9q34 and later on mutations in the *SETX* gene were identified^[71]. *SETX* is a large gene of 26 exons and encodes senataxin, a nuclear protein, which contains at its C terminus a classic seven-motif domain found in the superfamily 1 of helicases. The protein is involved in the response to oxidative stress^[53]. Acting as an RNA/DNA helicase, senataxin might have a significant role in the DNA repair pathway, in the splicing machinery^[71] and also in the processing of noncoding RNAs^[52]. Even though the *SETX* gene cannot be easily analysed by routine sequencing due to its size, several patients have been studied with this method and more than 75 different mutations have been identified thus far worldwide^[52,72,73] including missense, nonsense and splice site mutations, as well as small/gross deletions or insertions^[74].

(5) Ataxia with oculomotor apraxia type 3: Ataxia with oculomotor apraxia type 3 has more recently been described in a consanguineous Saudi Arabian family with four affected individuals that present with a similar phenotype to AOA2. Elevated serum α -fetoprotein levels were identified in all the patients and 2 siblings had OMA. The homozygous missense c.1885C > T [p.Pro629Ser] mutation in exon 12 of the *PIK3R5* gene has been associated with the disease in this family^[54]. *PIK3R5* is located on chromosome 17p12-p13. The encoded protein is a p101 regulatory subunit that interacts with class 1B PI3K which is involved in cell survival and growth as well as in metabolism, immune and cardiac functions. Even though it is registered as rs61761068 in db-single-nucleotide polymorphism (dbSNP), frequencies of the mutant allele in other populations indicate that it has only been detected in a heterozygous state, most likely representing non-affected carriers of the mutation^[54].

Spinocerebellar ataxia with axonal neuropathy: Spinocerebellar ataxia with axonal neuropathy is a related syndrome to the AOA group. It is characterized by the co-occurrence of cerebellar ataxia and axonal neuropathy. A single locus on chromosome 14q31-q32 has been associated with the disease and a missense mutation (c.1478A > G [p.His493Arg]) in the *TDP1* gene has been described in a large Saudi Arabian family^[75]. The encoded protein

tyrosyl-DNA phosphodiesterase 1 is a member of the DNA single-strand break repair complex and plays a significant role in the DNA repair pathway^[75,76].

Congenital and metabolic ataxias

Other ARCA are classified under the subgroups of congenital and metabolic ataxias (Table 1). Associated genes and mutations have been identified and molecular genetic testing by targeted mutation analysis or direct sequencing of the appropriate gene may be performed in ARCA patients based on the clinical picture. Moreover, in some cases of metabolic ataxias, biochemical analysis on specific available markers could precede and orientate the genetic testing.

Congenital ataxias: Congenital ataxias that have been associated with specific chromosomal loci and genes include the Joubert syndrome and the Cayman ataxia. Joubert syndrome has thus far been associated with seven loci (JBTS1, JBTS2, JBTS3, JBTS4, JBTS5, JBTS6, and JBTS7) and mutations have been identified in the seven *INPP5E*^[77], *TMEM216*^[78], *AHI1*^[79], *NPHP1*^[80], *CEP290*^[81], *TMEM67*^[82] and *RPGRIPL*^[83] genes respectively (Table 1). The Cayman ataxia has been described in an isolated population of the Cayman Island. A single locus on chromosome 19p13.3 has been associated with the disease and causative mutations in the *ATCAY* gene have thus far been identified^[84].

Metabolic ataxias: The subgroup of metabolic ataxias includes disorders characterized by progressive ataxia, recurrent ataxia and ataxia as a minor feature. Progressive ataxia includes ataxia with isolated vitamin E deficiency, abetalipoproteinemia, cerebrotendinous xanthomatosis and Refsum disease^[1]. Intermittent or minor ataxia includes: metachromatic leukodystrophy (*ARSA* gene)^[85,86], Niemann-Pick type C 1 (*NPC1* gene)^[87], GM1 (*GLB1* gene)^[88,89] and GM2 (*HEXA* gene)^[90,91] gangliosidosis, Wilson's disease (*ATP7B* gene)^[92-94], aceruloplasminemia (*CP* gene)^[95] and Chorea-acanthocytosis (*VPS13A* gene)^[96].

(1) Ataxia with isolated vitamin E deficiency: Ataxia with isolated vitamin E deficiency has a similar clinical picture with Friedreich ataxia. It is characterized by a biochemical abnormality, the very low plasma level of vitamin E and hence the low level of the α -tocopherol in the serum. The disease is caused by mutations in the α -TTP gene, encoding the α -tocopherol transfer protein which is responsible for the transfer of vitamin E to the nascent very low-density lipoproteins^[97]. Diagnosis may include the finding of low serum vitamin E values in the absence of malabsorption^[1] and further by performing genetic testing.

(2) Abetalipoproteinemia: Abetalipoproteinemia is characterized by deficiency of the liposoluble vitamin E, the low levels of cholesterol and the absence of the low-density lipoproteins. The clinical features comprise the

malabsorption syndrome, the pigmentary degeneration of the retina and progressive ataxic neuropathy. Moreover, patients show a characteristic deformation of their red cells called acanthocytosis^[98]. The disease is caused by mutations in the *MTP* gene which encodes the microsomal triglyceride transfer protein catalyzing the transport of triglyceride, cholesteryl ester and phospholipid between phospholipid surfaces^[99].

(3) Cerebrotendinous xanthomatosis: Cerebrotendinous xanthomatosis is a sterol storage disorder, characterized by abnormal accumulation of cholesterol and cholestanol in neural and other tissues, thus causing tendon xanthomas, premature atherosclerosis, cataracts and neurological dysfunction which is manifested as cerebellar ataxia, dementia and spinal cord paresis^[100]. The disease gene has been mapped to chromosome 2q33-qter and is caused by mutations in the *CYP27* gene have been identified. *CYP27* encodes sterol 27-hydroxylase, a mitochondrial cytochrome P450 enzyme that has an important role in cholesterol and bile acid metabolism^[100,101].

(4) Refsum's disease: Refsum's disease is caused by defective peroxisomal alpha-oxidation of phytanic acid, thus leading to its accumulation in blood and other tissues including neurons. It is characterized mainly by progressive retinitis pigmentosa, peripheral neuropathy and cerebellar ataxia. Other additional features include anosmia, deafness, cardiomyopathy, ichthyosis and skeletal abnormalities. The disorder is genetically heterogeneous since it has been mapped to two different loci. In most families, linkage was established to the first identified locus on chromosome 10pter-p11.2. Causative mutations have been found in the *PHYH* (or *PAHX*) gene^[102,103] encoding the phytanoyl-CoA hydroxylase enzyme that is involved in the catabolic pathway of phytanic acid. In a subset of patients mutations in the *PAHX* gene have been excluded. Further linkage analysis of these families led to mapping of a second locus on chromosome 6q22-24. In this subgroup of patients causative mutations have been identified in the *PEX7* gene encoding the peroxin 7 protein which is implicated in the import of matrix proteins into the peroxisome^[104].

Other known ARCA types

The following ARCA forms have been reported more recently that have not been included in the classification scheme described by Palau and Espinos (2006).

Spinocerebellar ataxia autosomal recessive type 8: Spinocerebellar ataxia autosomal recessive type 8 (SCAR8), also known as autosomal recessive cerebellar ataxia type 1 (ARCA1) is an adult-onset slowly progressive cerebellar ataxia accompanied by dysarthria and other associated features^[105]. It was first described and so far only reported in French-Canadian families most of them originating from defined regions of Quebec. A single locus on chromosome 6q25.1-q25.2 has been mapped and mutations

in the *SYNE1* gene have been associated with the disease. Syne-1 is involved in anchoring specialized myonuclei underneath the neuromuscular junction^[105] and plays a significant role in the development and maintenance of cerebellar functions. *SYNE1* is a large gene of 147 exons and this makes mutation screening by traditional methods expensive and complicated. Thus far direct sequencing of all the exons and flanking intronic sequences revealed a small number of missense and splice site mutations as well as a small scale deletion^[105,106].

Rundataxin-related ataxia: A new form of pure recessive cerebellar ataxia accompanied by epilepsy and mental retardation named 'Salih ataxia' has been described in a large consanguineous Saudi Arabian family with three affected individuals^[107]. Homozygosity mapping with SNP array (Gene Chip Human Mapping 10K 2.0 Xba Array, Affymetrix) and further linkage analysis with microsatellite markers, mapped the disease gene on chromosome 3q27.3-qter. Direct sequencing of the candidate genes enabled identification of the causative mutation in the *KIAA0226* gene. A single nucleotide deletion 2927delC [p.Ala943ValfsX146] frameshift mutation in this gene has thus far been identified. The encoded protein was named rundataxin based on its two conserved domains^[107] and it is associated with vesicular trafficking and signaling pathways.

NEXT-GENERATION SEQUENCING AND MOLECULAR DIAGNOSIS OF AUTOSOMAL RECESSIVE CEREBELLAR ATAXIAS

NGS technologies provide powerful tools for the identification of new genes. During the last 3 years, use of homozygosity mapping in combination with WES or targeted sequencing using the NGS platforms, enabled the identification of 5 new genes associated with rare types of ARCA in small families. WES has also been used as a screening tool thus enabling the rapid identification of mutations in known genes and demonstrating that it can be an efficient diagnostic tool.

Identification of new genes

Spinocerebellar ataxia autosomal recessive type 10: A new form of pure recessive ataxia accompanied by downbeat nystagmus and lower motor neuropathy, first described in a Dutch consanguineous family and named spinocerebellar ataxia autosomal recessive type 10 (SCAR10), was the first recessive ataxia molecularly diagnosed with homozygosity mapping and NGS^[108]. It is also known as ARCA3^[2]. Homozygosity mapping with a 10K SNP array (Affymetrix) combined with linkage studies with short tandem repeat markers led to fine mapping of the disease gene to chromosome 3p21.32-p22.3. A large number of genes were included in the mapped region thus making identification of the causative mutation

with the traditional Sanger sequencing difficult. Use of Sanger sequencing to analyze 15 candidate genes failed to identify any mutation. Targeted NGS was subsequently performed that enabled identification of the causative mutation in the *ANO10* gene, thus demonstrating the power of this method towards the identification of new genes. A homozygous c.1529T > G [p.Leu510Arg] mutation was identified in the Dutch family. Additional mutations have been identified in patients originating from Serbia (a homozygous frameshift mutation c.1150_1151del [p.Leu384fs]) and France (compound heterozygous mutations c.1476 + 1G > T and c.1604del [p.Leu535X]) through direct analysis of the new gene by conventional Sanger sequencing^[108]. More recently, a Japanese patient has been reported with a homozygous nonsense mutation (c.609C > G [p.Y203X]) identified by exome sequencing^[19] indicating that this new method can be applied as a screening tool allowing the fast identification of mutations in known genes. The encoded protein *ANO10* is a member of the human anoctamin family and similar to the other proteins of this family it is a putative calcium-dependent chloride channel, thus representing a new pathway implicated in cerebellar ataxias. However, this function remains to be confirmed^[108].

Spinocerebellar ataxia autosomal recessive type 11: A slowly progressive new type of ARCA accompanied by psychomotor retardation has been recently described in a small consanguineous Japanese family with two affected individuals and is known as spinocerebellar ataxia autosomal recessive type 11 (SCAR11). Homozygosity mapping using the Human SNP Array 6.0 (Affymetrix) combined with multipoint linkage analysis using extracted SNP array data revealed three candidate regions of homozygosity. WES was subsequently employed and data analysis focused on the regions of homozygosity. Sanger sequencing was used to further investigate filtered variants and the synaptotagmins 14 (SYT14) homozygous c.1451G > A [p.Gly484Asp] missense mutation has been associated with the disease. Further screening of all the coding regions of the entire gene in sporadic and familial ARCA cases failed to identify additional mutations. The encoded protein is Synaptotagmin XIV, a member of SYTs which are transmembrane proteins associated with exocytosis of secretory vesicles. Expression studies on SYT14 show that this protein plays a significant role in the cerebellum and support previous suggestions about the possible involvement of disrupted SYTs in neurodegeneration^[12].

Spinocerebellar ataxia autosomal recessive type 7: Spinocerebellar ataxia autosomal recessive type 7 (SCAR7) was first described in a Dutch family in 2004 as a distinct ARCA type, based on its genetic locus, the onset of the disease and/or other clinical findings^[109]. The disease gene has been mapped on chromosome 11p15 by genome-wide linkage analysis with microsatellite markers from the Marshfield genetic map. Identification of the causative mutation/gene was impossible for many years due to the large number (> 200) of genes within the

candidate region. Recently, the use of WES enabled the association of the disease in the family with two *TPP1* compound heterozygous mutations (the splice site c.509-1G > C and the missense c.1397T > G [p.Val466Gly])^[110]. These mutations were also detected in a sporadic SCAR7 patient. Mutations in the *TPP1* gene had already been associated with another disorder, the late infantile lipofuscinosis disease 2 (CLN2). Association of mutations in this gene with SCAR7 expands the phenotypes related to variants of the *TPP1* gene. Phenotype-genotype correlations proposed the hypothesis that CLN2 disease is due to variants causing the loss of enzyme activity, while SCAR7 is due to variants decreasing the enzyme activity. *TPP1* encodes the tripeptidyl-peptidase 1 enzyme, a member of the sedolisin family of serine proteases. This enzyme cleaves the N-terminal tripeptides from substrates and has a weak endopeptidase activity^[110].

Autosomal-recessive cerebellar ataxia with spasticity: A distinct form of ataxia previously described in three consanguineous Tunisian families with 7 affected individuals has been recently molecularly characterized with the combined strategy of homozygosity mapping and WES thus providing further evidence about the potential of these methods^[111]. Childhood or juvenile onset cerebellar ataxia was the main feature in all affected individuals of the families but later they developed pronounced limb spasticity. High density SNP genotyping was performed with the OmniExp-12, v1.0 DNA Analysis BeadChip (Illumina) and one large region of homozygosity on chromosome 9 was identified. The *APTX* gene mutated in AOA1 was included in the mapped region, so mutations in *APTX* were first excluded by Sanger sequencing. WES was then performed that revealed 2 causative mutations in the *GBA2* gene (the homozygous c.1018C > T [p.Arg340*] shared by two families and the homozygous c.2618G > A [p.Arg873His])^[111]. Sanger sequencing confirmed segregation of the identified mutations with the disease in the families. Furthermore, WES was used to screen 21 molecularly undiagnosed Tunisian individuals with ataxia. Through this procedure an additional *GBA2* mutation (c.363C > A [p.Tyr121*]) was identified^[111]. A second group associated 4 *GBA2* mutations with the spastic paraplegia type 46 (SPG46) phenotype^[112].

GBA2 encodes the non-lysosomal β -glucosidase 2 protein that was first recognized as a microsomal bile-acid β -glucosidase, while more recently it was reassigned to sphingolipid metabolism^[113], a critical process for the nervous system^[114]. The role of the enzyme in the central nervous system development was confirmed through the validation of the functional phenotype of some of the identified mutations causing HSP *in vivo* thus demonstrating the connection of sphingolipid metabolism and neurodegeneration^[112]. HSPs are another heterogeneous group of neurodegenerative disorders having common features with ataxias and as a consequence, many times it is difficult for the clinician to decide whether the patient has hereditary ataxia with spasticity or HSP with cerebellar features. Identification of mutations in the same gene

causing these two diseases demonstrates that a common pathway may be involved in neurodegeneration.

Spinocerebellar ataxia autosomal recessive type 13:

A new form of congenital cerebellar ataxia identified in Roma patients has recently been described and molecularly characterized^[113]. This new form is known as spinocerebellar ataxia autosomal recessive type 13 (SCAR13). WES was used both for linkage analysis and mutation detection. A subgroup of SNP genotypes was first selected from the exome sequence data and used to perform genome wide linkage analysis, thus emphasizing an additional application of this method. Linkage analysis mapped the disease gene to a single locus on chromosome 6q24. Data analysis was focused on the linked interval and finally resulted to the identification of two homozygous mutations in the *GRM1* gene (c.2652_2654del [p.Asn885del] and c.2660+2T > G). Sanger sequencing was used to verify the presence of mutations and their co-segregation with the disease within the families while PCR-based restriction-fragment-length-polymorphism assays and pyrosequencing were used to evaluate the mutation frequencies among control individuals. Both mutations interfere with a critical alternative splicing gene region. *GRM1* encodes the metabotropic glutamate receptor mGluR1 which is implicated in the modulation of intracellular Ca^{2+} levels and neuronal excitability. It is important for the development of the cerebellar cortex and the cerebellar and hippocampal synaptic plasticity, memory and learning^[115].

Identification of mutations in known genes

The combined strategy of homozygosity mapping and WES has been recently applied for the investigation of a consanguineous Turkish family and revealed a novel homozygous missense mutation (c.1366C > G [p.Leu456Val]) in the *C10orf2* gene, mutations in which have already been associated with infantile-onset spinocerebellar ataxia (IOSCA)^[16]. This rare type of ataxia was previously reported only in Finland. The recent identification of this novel mutation shows that IOSCA is not restricted to Finland.

WES was applied in another recent study to investigate two affected non-consanguineous siblings with a multisystem neurological disorder of unknown genetic origin and revealed two novel heterozygous mutations in the known *SACS* gene. These patients had been previously investigated with conventional methods which failed to identify the associated mutations in contrast to this novel approach which rapidly led to molecular diagnosis. Both patients shared the compound heterozygous mutations c.2076delG [p.Thr692Thrfs*713] and c.3965_3966delAC [p.Gly1322Valfs*1343] in the *SACS* gene that have been associated with ARSACS. Another novel homozygous mutation in *SACS* (c.2439-2440delAT [p.Val815Glyfs*4]) has been identified in a Tunisian family with two affected individuals, by homozygosity mapping and WES^[13]. A more recent study on a 4-year-old girl revealed an additional novel frameshift homozygous

mutation in the same gene by WES^[18].

WES also enabled the molecular diagnosis of the disease in an early onset spastic ataxia-neuropathy syndrome family. A homozygous missense mutation in the *AFG3L2* gene (c.1847G > A [p.Y616C]) has been identified^[116]. Heterozygous mutations in the *AFG3L2* gene have previously been identified in autosomal dominant spinocerebellar ataxia type 28 patients^[117,118]. The *AFG3L2* gene encodes a subunit of the m-AAA class of mitochondrial proteases which forms either a homooligomeric isoenzyme or a hetero-oligomeric complex with paraplegin, encoded by the *SPG7* gene^[116]. Mutations in the *SPG7* gene have been associated with hereditary SPG7, characterized by adult-onset spasticity and weakness of the lower extremities^[119]. In contrast to SCA28, this new syndrome associated with a recessive pattern of inheritance, is characterized by a much earlier onset and combines phenotypic features of SCA28 and *SPG7*^[116].

SYNOPSIS

The diagnosis of ARCA is becoming progressively more complex and challenging since clinical and genetic heterogeneity is expanding. Mutations in more than 20 genes have been associated with ARCA through the last decade, some of them identified in single families. There exists a significant number of patients remaining without a molecular diagnosis thus leading to the hypotheses that many other novel ARCA genes remain to be discovered in the future and that a different gene might be associated with the disease in each family mainly in populations with frequent inbreeding. The development of NGS technologies has already contributed significantly towards the molecular diagnosis of these rare diseases and appears very promising towards the diagnosis of additional cases in the future.

Sequencing of known candidate genes with conventional methods based on clinical findings has proved to be time-consuming and costly, except in the case of metabolic ataxias for which biochemical test results can precisely guide the molecular diagnosis. Routine investigation of some already known genes such as *SYNE1*, *ATM*, *SETX*, *POLG* and *SACS* is prohibitive, due to their significantly large coding regions.

Homozygosity mapping in consanguineous families has proved to be a powerful tool and may precede sequencing thus contributing to a more targeted analysis and facilitating the identification of novel ARCA genes. High density genome wide SNP genotyping with SNP arrays is the most rapid and effective method applied in the last years for homozygosity mapping. However, further analysis with more informative markers is often required in order to confirm or exclude some of the identified regions. Moreover, instead of using a SNP array, extracted SNP genotypes from the WES data may be used for genome wide linkage analysis. If a small number of genes exists within a mapped interval, then investigation by conventional sequencing may be efficient. Otherwise, if a large number of candidate genes exists, conventional

sequencing proves to be difficult, time consuming and expensive compared to the new methodologies.

WES enables the rapid sequencing of all coding regions of the genome and its usefulness has already been demonstrated in research and diagnostic applications. Conventional sequencing remains useful as a complementary method for the investigation of the segregation pattern of candidate variants previously revealed by WES and verification of the causative mutations. Moreover, investigation of families for mutations in a known gene within a mapped region could be performed either by conventional sequencing or by WES depending on the size of the coding regions that need to be examined. For small coding regions, the first method may be more cost effective while for large genes, WES is less expensive since costs have significantly decreased over the last year^[14].

Although the application of NGS technologies enables the identification of novel ARCA genes and the characterisation of novel mutations in known genes thus providing molecular diagnosis in many cases, there are some limitations due to their current reduced ability to sequence repetitive DNA expansions^[14]. The most common form of ARCA is caused by the GAA repeat expansion in intron 1 of the *FXN* gene. Additional forms of ARCA with unknown genetic etiology may also be associated with a similar type of mutation and will not be identified with the NGS approach.

CONCLUSION

Novel high-throughput sequencing technologies such as WES/WGS are ideal for simultaneously sequencing groups of candidate genes, whole-genome coding regions or the whole-genome and have brought a revolution in the molecular diagnosis of human genetic diseases. These technologies are capable of producing large amounts of data in one experiment and have been successfully used for the diagnosis of many rare and complex diseases including rare neurodegenerative diseases such as the ataxias. Their use for both research and diagnostic purposes has already contributed to the progress observed during the last years in the field of ataxias, including both the recessive and dominant forms. They are also very promising for the identification of many other new forms of ataxia in the near future since this group of disorders is very heterogeneous and a significant number of patients have already been excluded from many of the known genes. Determination of the genetic cause of each form is essential in order to initiate further investigation regarding the pathogenetic mechanisms and the potential therapies for these diseases.

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Pure motor stroke as the most frequent lacunar syndrome: A clinical update

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Core tip: Pure motor stroke (PMS) is the most common of any lacunar form (between one half and two thirds of cases). The posterior limb of the internal capsule, corona radiata, and pons are the most frequent brain topographies. This present update is focused on the clinical evidence and mechanisms underlying the relationship between PMS and different stroke etiologies.

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CLINICAL DEFINITION

Pure motor stroke (PMS), also known as pure motor hemiparesis, was first reported by Fisher and Curry in 1965 and it is considered the commonest lacunar syndrome in clinical practice, accounting for between one half and two thirds of cases, depending on the series. Lacunar infarction accounts for one fourth of all cerebral infarctions^[1,2] and generally develops in patients with hypertension and/or diabetes mellitus.

Definition of known pre-stroke hypertension includes patients having systolic and diastolic blood pressure higher than 140/90 mmHg, respectively, measured at least twice by the general practitioner before stroke onset, or those with high blood pressure at the time of stroke onset with one or more of the following: hypertensive retinopathy, left ventricular hypertrophy diagnosed by electrocardiographic or echocardiographic criteria, or abnormal renal function (excluding diabetes or other alternative cause of nephropathy). In accordance with World Health Organization criteria, diabetes is diagnosed

Abstract

Pure motor stroke (PMS), also known as pure motor hemiparesis, is the most common of any lacunar form (between one half and two thirds of cases, depending on the series). In an acute stroke registry, 733 patients presented a lacunar infarct and PMS accounted for 12.7% ($n = 342$) of all first-ever stroke patients and for 48% of all lacunar syndromes. The posterior limb of the internal capsule, corona radiata, and pons are the most frequent brain topographies. Infarcts in the mesencephalus or medullary pyramid have been exceptionally reported. This present update is focused on the clinical evidence and mechanisms underlying the relationship between PMS and different stroke etiologies.

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Table 1 Demographic characteristics and frequency of lacunar syndromes in the Sagrat Cor Hospital of Barcelona Stroke Registry

	<i>n</i> (%)
Total patients	733
Male	423 (57.7)
Age, yr, mean (SD)	74.1 (10.2)
Age ≥ 85 years old	110 (15.0)
Lacunar syndromes	
PMS	352 (48)
Pure sensory stroke	127 (17.4)
Sensorimotor stroke	83 (11.3)
Dysarthria-clumsy hand	59 (8)
Ataxic-hemiparesis	24 (3.3)
Atypical lacunar syndromes	88 (12)

PMS: Pure motor stroke.

Table 3 Topographies in pure motor stroke

Internal capsule
Pons
Corona radiata/centrum semiovale
Basal ganglia
Mesencephalon
Medullary pyramid
Cerebral cortex ¹

¹Cerebral topography related to PMS not due to lacunar infarct. PMS: Pure motor stroke.

when post-stroke repeated fasting plasma glucose levels exceeded 7.8 mmol/L (149 mg/dL). Serum determination of HbA1c is useful in doubtful cases to diagnose previous diabetes.

Pierre Marie in 1901 and Ferrand in 1902 had first stressed the link between isolated sudden hemiparesis and lacunes.

In the Sagrat Cor Hospital of Barcelona Stroke Registry, PMS accounted for 48% of all lacunar syndromes^[3] (Table 1). It was the first lacunar syndrome recognized clinically, and its features have been the most thoroughly explored^[1]. The clinical criteria of PMS include the presence of unilateral partial or complete paresis involving at least two of three areas (face, upper limb, or lower limb) of the body and no evidence of aphasia, apraxia, and agnosia, nor visual field defect, eye movement disturbance, ataxia, sensory loss or evidence of bilateral weakness^[4-6]. Pure motor monoparesis is almost never due to lacunar infarct (Table 2). These restricted motor deficits are more likely to be of cortical origin^[2,6].

Although the main elements of the PMS syndromes are motor, other complaints are not rare, especially sensory disturbances, which occur initially in as many as 42% of cases^[6]. These complaints usually present as numbness, heaviness, and loss of feeling. Only scan abnormalities are found on clinical examination. Improvement is seen in a high percentage of cases of PMS. Recovery is usually more rapid and more complete than that following cerebral surface infarction with a similar initial motor deficit^[7]. The syndromes of partial hemiparesis show

Table 2 Distribution of weakness related to pure motor stroke in 128 patients included in the Sagrat Cor Hospital of Barcelona Stroke Registry

Distribution of weakness	PMS (<i>n</i> = 128) <i>n</i> (%)
Face, arm, leg	112 (76)
Face, arm	6 (4)
Arm, leg	16 (10)
Face ¹	6 (4)
Arm	4 (3)
Leg	4 (3)

¹Patients with isolated pure facial central palsy with dysarthria have been included in the atypical lacunar syndrome in another study. Modified from Arboix *et al*^[6]. PMS: Pure motor stroke.

the best prognosis, as do those with the smaller infarct size on computed tomography scan or brain magnetic resonance imaging, but cases of complete hemiplegia followed by almost full recovery have been encountered^[5,8].

Lacunar infarctions are not a benign vascular condition; in contrast, they present a high risk of recurrence and vascular dementia in the mid and long term^[9-11].

Recurrent lacunar ischemic stroke may be associated with a more severe clinical picture and it is one of the major factors involved in producing lacunar state and vascular subcortical dementia^[7-12]. Recurrent strokes are more likely to be lacunar if the index event is lacunar^[5].

CLINICO-ANATOMIC CORRELATIONS

PMS has been reported from autopsied cases as a focal small (< 20 mm in diameter), deep infarction (lacunar infarct subtype) involving the internal capsule, corona radiata, pons and medullary pyramid^[1,2]. Other topographies are centrum semiovale, basal ganglia and mesencephalon (Table 3). The most common correlations have been with capsular locations. The greater number of lacunes has been reported in the posterior limb of internal capsule^[5].

These anatomical structures are supplied by the lentostriate branches and the long deep perforator medullary branches of the anterior and middle cerebral arteries, and the paramedian branches of the basilar artery^[1].

Small vessel diseases are mainly of atherosclerosis-degenerative type, such as lipohyalinosis or microatheromatosis of lacunar infarcts^[11,2].

OTHER CAUSES OF PURE MOTOR SYNDROMES

The concept of lacunar hypothesis (the presence of a lacunar syndrome is usually due to a lacunar infarct) suggested by Miller Fisher is clinically valid and useful in PMS^[13-15]. This lacunar syndrome is commonly due to lacunar infarct resulting from small vessel disease. In a clinical study^[5] of 222 patients with PMS, lacunar infarcts were found in 189 (85%) patients, whereas ischemic lacunar syndromes not due to lacunar infarcts occurred in 23 (10.4%) (atherothrombotic stroke in 12, cardioembolic

Table 4 Pure motor stroke not due to lacunar infarcts

Non-lacunar ischemic stroke ¹
Cardioembolic stroke
Atherothrombotic infarct
Internal carotid artery occlusion in the neck
Ventromedial pontine infarction due to a propagating thrombosis of the basilar branch
Complex aortic atheroma plaques
Ischemic stroke of unusual etiology
Ischemia-edema after craniotomy for postoperative bleeding
Intracerebral hemorrhage
Subdural hematoma
Brain abscess of the motor cortex
Cerebral metastasis
Demyelinating disease

¹Cerebral cortical surface infarction or large deep infarction (> 20 mm diameter).

stroke in 7, infarction of undetermined origin in 3 and infarction of unusual etiology in one) and hemorrhagic lacunar syndromes in 10 (4.5%). Other causes of PMS^[14,16] are showed in Table 4. These rare causes can nowadays be easily detected by imaging techniques.

However, the lacunar syndromes do not predict lacunar infarcts when used in the first hours of stroke. Later, they have a reasonable predictive power if the classic definition is retained^[17].

PURE MOTOR STROKE OF UNUSUAL ETIOLOGY

PMS and the other lacunar strokes may be caused in less than 5% of cases by other etiologies, mainly hematological diseases and infectious or inflammatory arteritis^[4,5,18].

Hematological disorders associated with lacunar stroke include polycythemia vera, essential thrombocythemia, and primary antiphospholipid antibody syndrome^[18].

Infectious arteritis due to neurosyphilis, neurocysticercosis, neuroborreliosis, and acquired immunodeficiency syndrome has been also associated with lacunar infarction. In patients with first ever stroke, chronic *Helicobacter pylori* infection detected by IgG antibodies is associated with risk of small artery occlusion^[5].

Moreover, inflammatory arteritis in systemic lupus erythematosus or granulomatous angiitis and polyarteritis nodosa have been related to lacunar infarctions, although in cases of polyarteritis nodosa, lacunar strokes are due to thrombotic microangiopathy, not vasculitis^[18].

Drug abuse, particularly of cocaine, and pseudoxanthoma elasticum may produce small, deep infarcts^[5].

Other uncommon etiologies include hereditary (cerebral autosomal dominant arteriopathy with subcortical infarcts and leukoencephalopathy), mitochondrial (mitochondrial encephalopathy, lactic acidosis, and stroke-like episodes), deposition of substances in the vessel wall (amyloid in sporadic and hereditary cerebral amyloid angiopathy or ceramide in Fabry's disease), or toxic cause. Other small vessel diseases are venous collageno-

sis, post-radiation angiopathy and non-amyloid microvessel degeneration in Alzheimer disease^[5,18].

CONCLUSION

PMS is the most common lacunar syndrome. The lacunar hypothesis is clinically valid and useful in PMS; however in 10%-15% of the cases PMS may be associated with underlying non-lacunar ischemic mechanisms such as cardiac source of embolism, atherosclerosis, intracerebral hemorrhage or cerebral ischemia of unusual etiology that may influence management. Thus, it is mandatory to establish the etiological diagnosis of PMS to classify correctly the stroke subtype.

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Overview of botulinum toxin as a treatment for spasticity in stroke patients

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Core tip: In this overview, we describe that botulinum toxins (BTXs) were very useful to reduce spasticity in stroke patients. Moreover, we introduce studies on appropriate effective and safe injection methods; pilot studies that evaluated combined rehabilitation and BTX treatment; and adverse events after BTX injection, including risk factors of neutralizing antibodies.

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Abstract

Spasticity after the occurrence of stroke induces limb deformity, functional disability and/or pain in patients, which limits their activities of daily living and deteriorates their quality of life. Botulinum toxin (BTX) has recently been reported as an efficacious therapeutic agent for the treatment of spasticity. Systematic review and meta-analysis studies have demonstrated that BTX therapy after stroke reduces spasticity and increases physical activity capacity and performance levels. Moreover, BTX can be used as an adjuvant in physiotherapy. Several studies have confirmed that the combination of BTX therapy and physiotherapy improves motor recovery. However, to date, only a few such combination studies have been conducted and their findings are considered preliminary and controversial. Therefore, future studies are required to determine the appropriate combination of treatment methods that will aid motor recovery.

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Key words: Botulinum toxin; Rehabilitation; Stroke;

INTRODUCTION

Spasticity is a common complication in patients who have experienced stroke. The incidence of spasticity is reported to be 17%-43% in post-stroke patients^[1-3]. Spasticity induces limb deformity, functional disability and/or pain that limits activities of daily living (ADL) and deteriorates the quality of life (QOL). In order to reduce spasticity, clinicians prescribe oral medications, intrathecal baclofen therapy, nerve phenolization, casting or splinting, rehabilitation therapy or a combination of these therapeutic approaches^[4-8]. A recent study has shown that botulinum toxin (BTX) is an efficacious therapeutic agent for the treatment of spasticity^[9].

BTX is poisonous in nature and is produced by the anaerobic bacterium *Clostridium Botulinum*^[10]. BTX blocks vesicular acetylcholine release at neuromuscular junctions and reduces focal muscle tonus and contracture^[11]. Therefore, BTX is used for treating muscular hyperactivity and excessive or inappropriate muscle contraction. Seven types of BTX exist in nature, but two toxin types, type A (BTX-A) and type B (BTX-B), are used in the clinical

setting. Most clinical trials have utilized BTX-A because it has a longer lasting effect than BTX-B^[12,13]. BTX-B tends to be selected for spasticity treatment when patients have neutralizing antibodies against BTX-A or develop antibodies after repetitive BTX-A injection treatment^[14,15].

Many studies have reported that BTX injection treatment reduces spasticity in stroke patients^[16-25]. However, a few studies have focused on rehabilitation therapy after BTX treatment for improving motor function^[26-28]. This review focuses on the following factors of BTX treatment in post-stroke patients: (1) the effectiveness of spasticity reduction, with regard to ADL and QOL; (2) the appropriate injection method; (3) combined BTX therapy and neurorehabilitation; and (4) adverse events. The purpose of this review is to provide a comprehensive overview of BTX treatment for muscular spasticity in post-stroke patients, to understand its mechanisms of action, and to suggest appropriate treatment approaches.

EFFECTIVENESS FOR SPASTICITY

Several reports clarified the effectiveness of BTX treatment for upper limb spasticity experienced after stroke^[16-21]. A recent systematic review and meta-analysis reported that BTX-A reduced spasticity and thereby improved physical activity capacity and performance levels in the upper limb in patients who experienced stroke^[29]. BTX-A (Botox[®]) injection in the upper limb muscles are recommended at the following doses: 25-100 units for the flexor carpi radialis; 20-70 units for the flexor carpi ulnaris; 20-60 units for the flexor digitorum superficialis; 20-60 units for the flexor digitorum profundus; 10-30 units for the flexor pollicis longus; and 5-25 units for the adductor pollicis^[30].

Furthermore, several studies have reported the effectiveness of BTX therapy in stroke-induced lower limb spasticity^[22-25]. However, a systematic review and meta-analysis reported that BTX-A treatment for stroke-induced lower limb spasticity was associated with a slight improvement in functional ambulation^[19,31]. A multicenter, double-blind trial demonstrated the effectiveness of BTX-A in cases of lower limb spasticity, but improvements in gait pattern scale and speed were not significant, compared to that observed in the placebo group^[32]. A previous study suggested that high-dose BTX-A treatment may induce excessive muscle weakening that may deteriorate limb function^[25]. BTX-A (Botox[®]) injection doses in the lower limb is recommended at the following doses: 50-250 units for the medial and lateral head of the gastrocnemius respectively; 50-200 units for the soleus; and 50-150 units for the tibialis posterior^[30]. However, to the best of our knowledge, no study has evaluated the appropriate BTX dose for treatment of upper and lower limb spasticity and therefore, future studies are required to clarify these dosages. Moreover, it is noted that clinicians should consider the maximum dose of BTX per session, which differs in each country (*e.g.*, 360 units in Japan and 600 units in Europe)^[32-35].

Treatment of spasticity in patients who experience a

stroke with BTX-A may improve ADL and QOL^[36,37]. A previous study reported that high-dose BTX-A significantly improved ADL scores for limb position and self-dressing^[36]. Furthermore, another study in post-stroke patients reported that BTX-A therapy improved overall QOL scores and muscle tone, reduced disability, and improved the ability to function^[37]. However, a recent meta-analysis study revealed that BTX-A did not significantly improve overall health-related QOL^[38].

APPROPRIATE INJECTION METHOD

Several studies have reported on BTX-A injection techniques^[39-41]. Francisco *et al.*^[39] compared the efficacy of high and low volumes of BTX at the same dose for wrist and finger flexor spasticity. They found that modified Ashworth Scale scores were significantly decreased in both high and low-volume injection groups. However, there was no significant difference between the groups^[40]. Gracies *et al.*^[41] reported that injecting a small volume of BTX-A near the neuromuscular junction was more effective than injecting a similar volume at a greater distance from the neuromuscular junction. However, injecting a large volume at a greater distance from the junction was as effective as a small volume injected near the neuromuscular junction. Mayer *et al.*^[40] also reported that a single motor point and multisite low-dose, high-volume BTX-A injections produced a similar impact in spasticity reduction. Therefore, the appropriate injection methods according to BTX-A volume and injection points must be elucidated. Moreover, BTX-A injections are more accurate when the affected muscles are targeted *via* needle electromyography or under ultrasound guidance. This conventional method is not supported by evidence, but correct muscle selection has been confirmed to be a key feature in the efficacy of BTX treatment^[42,43].

COMBINATION OF BTX THERAPY AND NEUROREHABILITATION

As mentioned above, BTX improves motor function by reducing muscular spasticity in post-stroke patients. Therefore, BTX therapy may be a useful adjuvant therapy in combination with neurorehabilitation. In fact, several studies have reported that the combination of BTX therapy and physiotherapy improved motor recovery^[26-28]. A recent study reported that BTX injection, followed by home-based functional training, may have the potential to improve active motor function of the affected upper limb in post-stroke patients^[26]. Moreover, previous studies have reported that casting and dynamic splinting following BTX-A injection improved range of motion as well as motor function^[5,27,28]. Thus, BTX when combined with physiotherapy or casting may be useful for improving motor function in stroke-induced spasticity.

Recently, various neurorehabilitation strategies have been established to improve neural plasticity and motor recovery. These approaches include constraint-induced

movement therapy (CIMT), transcutaneous electrical neuromuscular stimulation (TENS) and non-invasive brain stimulation (NIBS)^[44-49]. A preliminary study demonstrated that the combination of BTX injection and CIMT may improve functional gains in post-stroke patients^[44,45]. Moreover, a recent study reported that combined BTX-A and NIBS improves motor function in patients with stroke-induced upper limb hemiparesis^[48]. However, a previous study reported that combining BTX injection with TENS did not have any additional effect on motor recovery^[47]. The number of these combination studies remains small and their results are considered preliminary and controversial. Although BTX-A probably enhances the effects of neurorehabilitation strategies in post-stroke patients, future studies are required to determine the appropriate combination of treatments for optimum motor recovery.

ADVERSE EVENTS

In general, pain, rash, edema and muscle weakness have been noted at the injection site and remote muscle weakness, fatigue and allergic reaction have been reported as adverse events related to BTX therapy. To clarify the safety of BTX-A, Turkel *et al.*^[50] analyzed 9 double-blind, placebo-controlled studies in more than 500 patients who had experienced a stroke. The most frequent reported adverse events by the patients (> 5% but < 10% in either group) were respiratory infection, seizures, incoordination and injection site pain, none of which occurred at a significantly high rate in the BTX-A group. Nausea alone was reported at a significantly higher rate in the BTX-A group than in the placebo group. In contrast, injection site pain, chest pain and allergic reaction were found at a significantly higher frequency in the placebo group. Other reviews have also reported no serious adverse events after BTX-A administration in adult post-stroke patients^[35]. Therefore, BTX has recently been considered a safe and efficacious treatment for spasticity in post-stroke patients.

However, care should be taken to try to avoid producing neutralizing antibodies. A higher dose per treatment, repetitive treatment and shorter intervals between treatments are risk factors for the production of neutralizing antibodies^[51-53]. Of these, the interval between treatments is the most important risk factor. Therefore, BTX-A treatment intervals of more than three months and BTX-B treatment intervals of two months are recommended^[54].

CONCLUSION

Many reports have shown the usefulness of the BTX treatment for spasticity in stroke patients. However, a few studies have reported the effectiveness of rehabilitation after BTX treatment in stroke patients. Future studies are needed to evaluate the efficacy of BTX after treatment with rehabilitation.

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Neuritin: A therapeutic candidate for promoting axonal regeneration

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Abstract

Following injury, the axons of the mammalian central nervous system do not regenerate. Many studies have aimed at understanding the mechanisms that prevent axonal regeneration and at designing ways to overcome the obstacles preventing axonal regrowth. These studies have identified numerous proteins as promoters of axonal regeneration. In this minireviews, we focus on neuritin as a therapeutic candidate for promoting axonal regeneration. Neuritin was first identified as a neuronal-activity-inducible gene product in the rat brain. The overexpression of neuritin in neurons or the application of neuritin to neurons induces neuritogenesis, neurite arborization, and axonal elongation both *in vitro* and *in vivo*. These morphological changes are often observed during the first step of axonal regeneration. Indeed, neuritin expression increases during axonal regeneration in the peripheral nervous system (PNS). Conversely, in a mouse model of diabetes mellitus, neuritin expression decreases in the PNS, and this reduced expression may result in deficient axonal regeneration. Neuritin is induced in the hippocampal dentate gyrus after temporal lobe epilepsy or brain ischemia; however, in these conditions, neuritin induc-

tion may exacerbate brain dysfunction through mossy fiber sprouting. Together, these findings support the hypothesis that tightly controlled regulation of neuritin may be required for the treatment of each unique axonal pathology.

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Key words: Axonal regeneration; Neuronal development; Axonal injury; Neuritogenesis; Epilepsy

Core tip: Neuritin has been shown to be an activity-regulated protein in neurons. Its expression also increases after neuronal damage. Neuritin induces neuritogenesis, arborization, and axonal elongation. These functions may be beneficial for axonal regeneration after nerve injury. Here, we review neuritin as a therapeutic candidate for promoting axonal regeneration in the peripheral and central nervous systems. We also discuss the possible involvement of neuritin in mossy fiber sprouting after epileptic seizures or brain ischemia.

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INTRODUCTION

After neuronal injury, axonal regeneration is not observed in the mammalian central nervous system (CNS). The regeneration of an axon itself is the only way to recover neuronal function after CNS injury. For decades, many investigators have tried to develop methods that promote axonal regeneration.

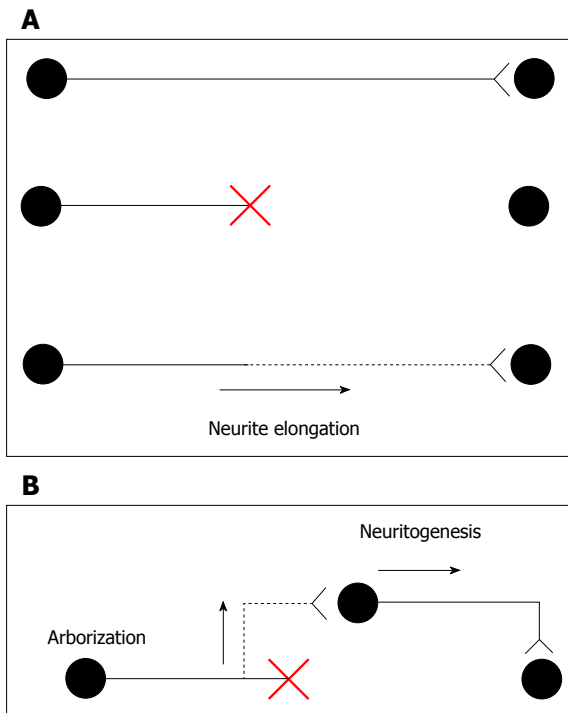


Figure 1 Schematic illustration of axonal regeneration. A: Top, intact axon; middle, transected or crushed axon; bottom, canonical axon regeneration. New growth occurs from the tip of the transected axon, and the regenerating axon reinnervates its normal target; B: Regenerating axon: the branch arises from the axon close to the injury site. The new axon branch connects to the surrounding neuron, which extends an axon to the original target. Figures are modified from Tuszynski *et al.*^[60].

In canonical axonal regeneration, new growth is expected to occur from the tip of a transected axon and reinnervate its normal target. Because CNS neurons maintain the intrinsic ability of axonal outgrowth^[1], the promotion of neurite outgrowth can contribute to the regrowth of axons (Figure 1A). In addition to canonical regeneration, other processes can achieve the functional recovery of transected neurons. New axon branches could arise from the axonal region close to the injury site, and these branches could connect with other newborn neurons near the lesion. Alternatively, neural precursor cell grafts enable the production of newborn neurons. New neurons close to the injury site are expected to extend their axons to the target organs. In this model, enhancements of neuritogenesis and arborization help form new axons and branches, respectively (Figure 1B). A few transplant models have succeeded in promoting axonal functional recovery^[2,3]. Thus, axonal elongation, branch formation, and neuritogenesis may be crucial for axonal regeneration.

In this minireviews, we focus on the protein neuritin, which is one candidate for the promotion of axonal regeneration. The expression level of neuritin is altered in CNS and peripheral nervous system (PNS) diseases. In addition, the overexpression or application of neuritin promotes neurite formation, branch formation, and neurite elongation in cultured neurons. These changes may promote axonal regeneration.

NEURITIN: ITS CHARACTERISTICS AND DISTRIBUTION

Neuritin, which is also known as candidate plasticity gene 15 (CPG15), was first identified during a screen for neural-activity-regulated genes in the hippocampus^[4]. Neuritin mRNA is predominantly expressed in the brain. Minor expression is observed in the lungs and liver^[5]. Within the brain, the highest levels of expression are in the dentate gyrus of the hippocampus^[5]. Expression of neuritin is increased by neural stimulation. Neuritin mRNA was increased both in the cerebral cortex by light^[6], sensory experience^[7], and exercise^[8] and in the hippocampus by exercise^[8] and electroconvulsive seizure^[9]. Immunohistochemical analysis showed the expression of neuritin in the sciatic nerve^[10] and facial motor neuron^[11], indicating that neuritin is also expressed in the PNS.

An open reading frame of the neuritin sequence predicts a protein with 143 amino acids and a 27-amino-acid signal peptide. The 27 C-terminal amino acids contain a consensus cleavage signal that is found in glycosylphosphatidylinositol-anchored proteins. The predicted mature protein is a membrane-bound 12-kDa protein with no apparent phosphorylation or glycosylation site. Neuritin shows no significant homology with any known protein^[5]. A portion of neuritin may be cleaved from the membrane and exists as a soluble form^[12].

Neuritin protein is concentrated in the neuronal soma and is unevenly dispersed along neuritic projections in the rat brain^[5]. Neuritin is expressed in puncta throughout cell bodies, dendrites, and axons, although its expression in axon terminals is particularly strong^[13].

NEURONAL FUNCTIONS OF NEURITIN

The neuronal functions of neuritin have been investigated primarily through studies involving the overexpression, gene silencing, and application of neuritin protein. These functions of neuritin can be categorized into the following three groups: neuritogenesis, neurite arborization, and neurite extension (Figure 2A).

Neuritogenesis

After the administration of purified neuritin to cultured hippocampal and cortical neurons, neurons showed extensive dose-dependent neuritogenesis compared with control neurons^[5,13]. In contrast, gene silencing of endogenous neuritin caused a robust reduction in nerve growth factor (NGF)-induced neurite formation^[14]. Because NGF induces neuritin expression in dorsal root ganglion (DRG) neurons^[10], NGF may promote neurite formation through neuritin expression; however, it remains unclear how NGF regulates neuritin expression. Furthermore, the application of a neutralizing antibody against neuritin to cultured cortical neurons significantly decreased the number of their primary dendrites and their total dendritic length^[13]. These findings suggest that neuritin enhances neuritogenesis in developing neurons.

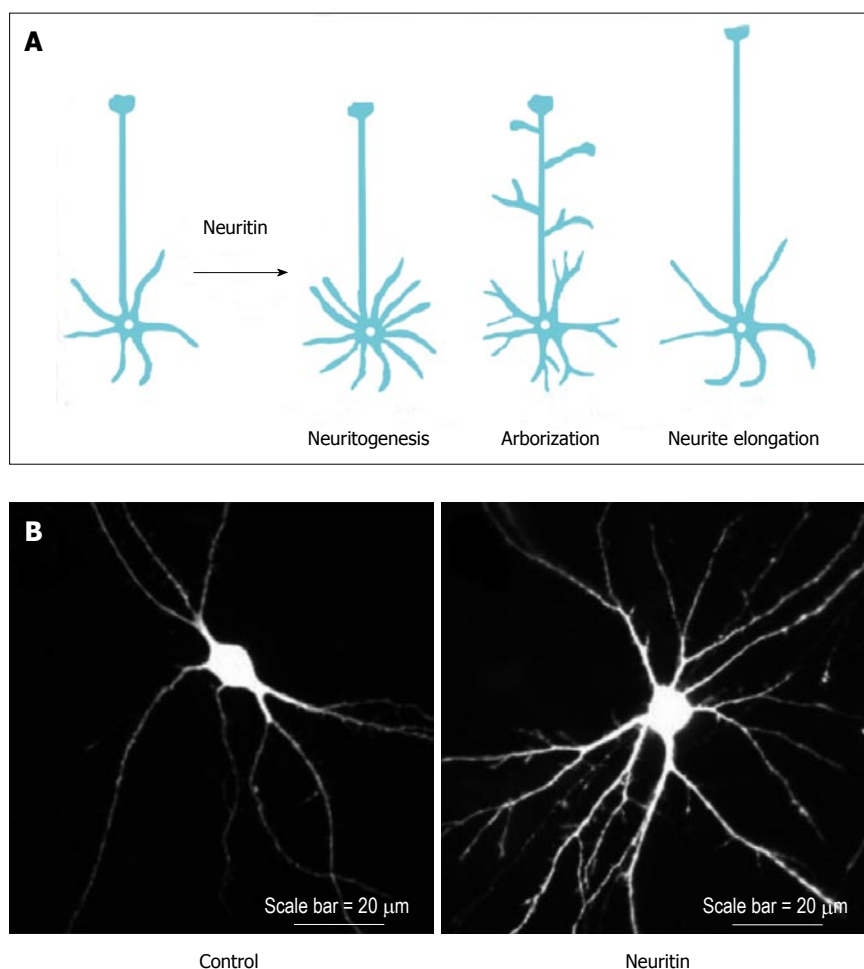


Figure 2 Neuritin induces neurite arborization. A: Neuritin regulates neuritogenesis, neurite arborization, and neurite extension; B: Hippocampal neurons were prepared from rat embryos that were transfected with green fluorescent protein (GFP) (left) or GFP and neuritin (right) after 5 d *in vitro*, and then they were immunostained after 12 d *in vitro*.

Neurite arborization

Neuritin can promote branch formation in neuronal dendrites. The overexpression of neuritin increased dendritic arbors in the optic tectal neurons of *Xenopus* tadpoles^[15]. Cultured rat hippocampal neurons overexpressing neuritin also exhibited significant increases in dendrite arborization (Figure 2B). Furthermore, neuritin targets axons to promote branch formation. The overexpression of neuritin changed the dynamic behavior of axonal branches. In the neuritin-expressing *Xenopus* motoneurons, the axonal branches grew significantly faster than those of control neurons owing to increased branch formation and decreased branch retraction^[16].

Neurite extension

In addition to neuritogenesis and branch formation, neuritin promotes the elongation of axons and dendrites. Dissociated cortical neurons plated on neuritin-coated dishes extended longer neurites than those plated on bovine serum albumin-coated dishes^[5]. In addition, dissected hippocampal explants co-cultured with neuritin-expressing cells showed significantly longer neurites than those co-cultured with control cells^[17]. Conversely, down regulation of neuritin expression completely abolished NGF-induced neurite outgrowth. The longest neurite of neuritin-reduced neurons treated with NGF was comparable to that of control neurons cultured without

NGF^[10]. Thus, there has been progress in our understanding of neuritin's functions, but by contrast, little is known about its mechanisms of action. A recent study has shed light on its signaling pathway (*e.g.*, the effects of neuritin on cerebellar granule neurons is attenuated by ERK, Akt, or mTOR inhibitors^[18]); however, further work will be needed to dissect the mechanistic contributions of neuritin to different forms of neuronal function.

POSSIBLE ROLES OF NEURITIN IN AXONAL REGENERATION

Neuritogenesis, axonal elongation, and axonal branching may be necessary for axonal regeneration. Accordingly, neuritin could contribute to axonal recovery. Several lines of evidence indicate that neuritin is involved in axonal regeneration in the PNS (Figure 3). In diabetes mellitus, axonal regeneration is nearly nonexistent in both experimental animal models^[19,20] and clinical cases^[21,22]. Streptozotocin (STZ) treatment causes experimental diabetic neuropathy in rats. Neuritin mRNA and protein have been shown to be significantly reduced in the DRG after STZ treatment^[10]. The sections proximal to a ligation in the sciatic nerve exhibited a significant reduction in neuritin levels in diabetic rats compared with those

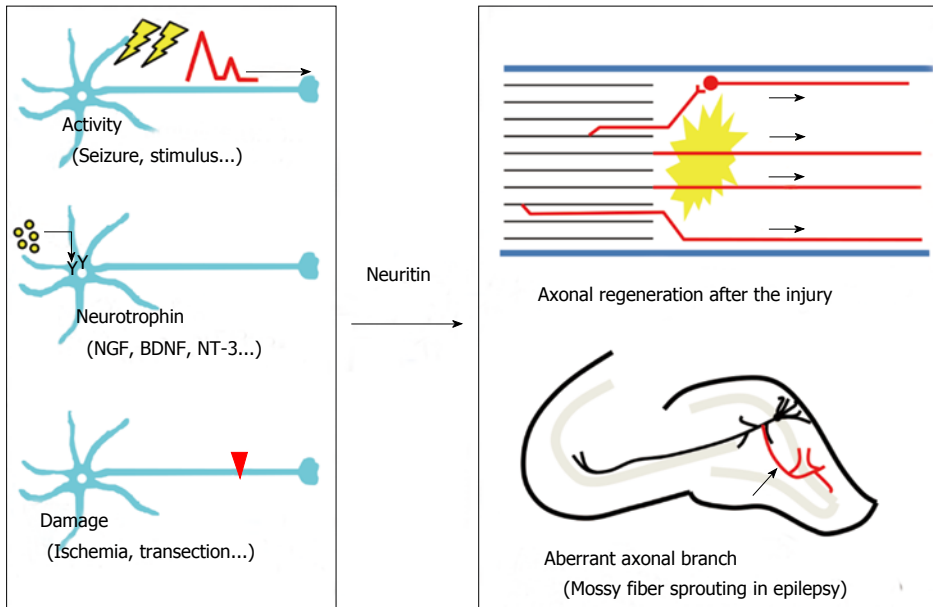


Figure 3 Induction of neuritin and its pathophysiological functions. Representative neuronal events that increase the expression of neuritin. Seizures, neurotrophins, and neuronal damage induce the expression of neuritin (left). The physiological roles of neuritin. Neuritin-induced changes might promote axonal regeneration after nerve injury, whereas the same morphological changes could exacerbate temporal lobe epilepsy (right). NGF: Nerve growth factor; BDNF: Brain-derived neurotrophic factor; NT-3: Neurotrophin-3.

in control rats, while distal sites did not^[10]. These results suggest that the axonal transportation of neuritin to the ligature site may be reduced in diabetic rats, raising the possibility that experimental diabetic neuropathy could result from a deficiency in axonal regeneration caused by a decrease in neuritin expression.

What mechanism is involved in the reduction of neuritin in diabetic neuropathy? In cultured sensory neurons, NGF has been shown to increase neuritin mRNA and protein in a dose-dependent manner. NGF treatment reverses the decrease in neuritin levels in the DRGs of STZ-induced diabetic rats^[10]. In addition, NGF levels are reduced in the diabetic PNS^[23,24]. These results indicate that exogenous NGF may promote axonal regeneration in human diabetic patients *via* neuritin induction.

Androgen has been shown to induce axonal regeneration after the axotomy of various motor neurons^[25]. After axotomy of the hamster facial nerve, exogenous androgen treatment lead to functional recovery by axonal regeneration^[26]. Androgen treatment after axotomy of the facial nerve increased the expression of neuritin mRNA *in vivo*^[11]. This androgen-dependent increase in neuritin mRNA was blocked by an antagonist of the androgen receptor^[27]. Androgen-induced neuritin expression may participate in the axonal regeneration of other motor neuron injuries. Androgen may induce neuritin expression through a different mechanism than NGF signaling does. The neuritin promoter harbors seven putative androgen response elements^[27], suggesting that the androgen receptor may directly control neuritin gene expression in motor neurons.

Neuritin mRNA has also been observed to increase after spinal cord injury. Fourteen days after injury by the

weight-drop method, the amount of neuritin mRNA peaked near the injury site^[28]. This finding may suggest that neuritin contributes to axonal regeneration in the CNS as well as in the PNS; however, future experiments are required to provide more definitive evidence. In addition, many questions remain regarding the molecular mechanisms by which neuritin expression is regulated under each pathological condition.

INVOLVEMENT OF NEURITIN IN EPILEPTOGENESIS

Neuritin was first identified as a neural-activity-regulated gene product in the brain. In addition to the induction following spinal cord injury, neuritin expression is also drastically increased in the hippocampus in models of epilepsy and brain ischemia (Figure 3).

In temporal lobe epilepsy, granule cells in the hippocampal dentate gyrus show axonal sprouting and branching. This axonal change, which is called mossy fiber sprouting, is one of the hallmarks of an epileptic hippocampus observed in both experimental animal models^[29-33] and clinical cases^[34-38]. Mossy fiber sprouting leads to axonal projections into the inner molecular layer of the dentate gyrus and results in the formation of a closed recurrent circuit^[29,39-42]. The formation of a new excitatory circuit within the dentate granule cells induces spontaneous bursting in the dentate gyrus under certain conditions^[33,43-45]. The idea that mossy fiber sprouting is the main reason for acquired epileptogenesis is controversial; however, it is probable that increased axonal branching in the granule cells results in the formation of epileptogenic recurrent circuits in the hippocampus^[46,47].

We have observed that neuritin promotes the arborization and elongation of mossy fibers in hippocampal slice cultures (unpublished data). This neuritin-induced mossy fiber sprouting may contribute to the formation of closed circuits between the granule cells (Figure 3). In this instance, an upregulation of neuritin may be harmful to brain function. Inhibition of neuritin function could be necessary for the suppression of epileptogenesis in temporal lobe epilepsy.

Transient middle cerebral artery occlusion (MCAO) leads to permanent damage of the brain that affects the lateral striatum and the lateral parietal, temporal and occipital cortex^[48]. Two hours after MCAO treatment in rats, an induction of neuritin mRNA was observed in the ipsilateral dentate gyrus of the hippocampus, as well as in the ipsilateral cerebral cortex^[49]. The increase in the hippocampus was robust and confined to this area 6 h after reperfusion^[49]. In this model, the significance of neuritin induction in the cortex remains unclear; however, the upregulation of neuritin in dentate granule cells might indicate its involvement in the pathogenesis of postischemic epilepsy through mossy fiber sprouting (Figure 3).

Thus, aberrant neuritogenesis, axonal branch formation, and axonal elongation by neuritin could exacerbate neuronal dysfunction in the CNS. Tightly controlled regulation of neuritin expression may be required for the treatment of various CNS diseases.

CONCLUSION

Neuritin is a small extracellular protein that is expressed in neurons, and its expression is upregulated by neuronal activity. The expression of neuritin increases in response to neuronal damage and neurotrophin treatment in the CNS and the PNS. Neuritin expression leads to morphological changes in neurons. The induction of neuritogenesis, the enhancement of neurite arborization, and the promotion of neurite extension are all elicited by neuritin. These morphological changes suggest that neuritin can contribute to axonal regeneration after axotomy. A variety of studies provide evidence that neuritin may contribute to axonal regeneration in PNS and CNS injuries. In contrast, neuritin induction in the hippocampus in response to epileptic seizures or brain ischemia may aggravate neuronal damage by enhancing mossy fiber sprouting. Thus, tightly controlled regulation of appropriate neuritin levels will be useful in the treatment of axonal pathology.

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Antiplatelet strategy for acute ischemic stroke: A mini review

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Core tip: Patients with acute ischemic stroke have a high risk of deterioration in the 48-72 h after symptom onset. Although thrombolysis is an effective method, most patients are excluded due to the limit of strict indications. Early antiplatelets are recommended for most patients. However, the question remains unclear whether another effective and safe antiplatelet strategy for the treatment of acute ischemic stroke exists. Growing evidence shows that combination antiplatelets may be superior to mono antiplatelets in the treatment of acute ischemic stroke.

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Abstract

Transient ischemic attacks and minor ischemic strokes have a high risk of an unstable clinical course in the initial 48-72 h after symptom onset. Early antiplatelet treatment is recommended to treat most patients with acute ischemic stroke because few patients can be treated with thrombolysis due to the limit of strict indications, such as a time window. Antiplatelets aim to prevent recurrence or deterioration of stroke. The guidelines recommend the use of aspirin in the acute stage based on two clinical trials. However, some patients still developed recurrence or deterioration of stroke despite timely aspirin administration. Thus, the question remains unclear whether another effective and safe antiplatelet strategy for the treatment of acute ischemic stroke exists. Growing evidence shows that combination antiplatelets may be superior to mono antiplatelets in the treatment of acute ischemic stroke.

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Key words: Antiplatelet; Acute ischemic stroke

INTRODUCTION

In the acute stage of ischemic stroke, one of the most important treatment strategies aims to prevent recurrence or deterioration of stroke, because patients with transient ischemic attack (TIA) and minor ischemic stroke have a high risk of an unstable clinical course. Most of these unfavorable events occur in the initial 48-72 h after symptom onset and have obvious effects on the recovery of neurological functions in these patients^[1-4]. It is well demonstrated that thrombolysis is an effective method to treat acute ischemic stroke. Due to the limit of strict indications, such as a time window, most patients are excluded for thrombolysis. Thus, early antiplatelet treatment is recommended to treat most patients with acute ischemic stroke in the guidelines^[5,6]. In acute ischemic stroke, 2 mega trials established the efficacy of aspirin^[7,8]; thus, the guidelines recommend the use of aspirin in the acute stage^[5,6]. However, some patients develop recurrence or deterioration of stroke despite timely aspirin administration. Thus, the question remains unclear whether another effective and safe

antiplatelet strategy for the treatment of acute ischemic stroke exists.

In this article, we will briefly review the following topics: (1) the action mechanisms of different antiplatelet agents; (2) the evidence for antiplatelet therapy for acute ischemic stroke; and (3) the efficacy and safety of combination antiplatelet agents for acute ischemic stroke.

ANTIPLATELET DRUGS AND MECHANISMS

Commonly used antiplatelet agents include aspirin, clopidogrel, cilostazol and dipyridamole. These agents act at different sites of the platelet aggregation pathway to inhibit platelet activation. Aspirin is a non-selective and irreversible cyclooxygenase (COX) inhibitor, which selectively acetylates the hydroxyl group of the COX enzyme, thus inhibiting conversion of arachidonate to prostaglandin G₂/H₂ and thromboxane A₂. This leads to irreversible inhibition of platelet aggregation^[9].

Clopidogrel is a thienopyridine derivative that prevents ADP-induced platelet activation and aggregation by irreversibly inhibiting the binding of adenosine 5-diphosphate to glycoprotein IIb/IIIa, with the result of preventing the binding of fibrinogen to this receptor^[10].

Cilostazol is an inhibitor of phosphodiesterase 3 (PDE3) which results in increased concentrations of intracellular cAMP^[11]. Cilostazol inhibits primary and secondary platelet aggregation and high-shear stress platelet aggregation both *in vivo* and *in vitro*^[12].

Dipyridamole is postulated to have multiple mechanisms of action. On one hand, its inhibition of platelet phosphodiesterase produces an increase in intraplatelet cyclic adenosine monophosphate level, resulting in the potentiation of the platelet inhibitory actions of prostacyclin. On the other hand, it also acts by directly releasing eicosanoid from vascular endothelium and inhibiting cellular uptake and metabolism of adenosine. This leads to the inhibition of platelet aggregation again^[13].

ANTIPLATELET THERAPY FOR ACUTE ISCHEMIC STROKE

A lot of studies have investigated the efficacy and safety of antiplatelet treatment in the secondary prevention of ischemic stroke^[14]; however, only a few studies have investigated the efficacy and safety of antiplatelet treatment in acute ischemic stroke^[7,8,15,16]. Of these studies, there are two powerful clinical trials: The Chinese Acute Stroke Trial (CAST)^[8] and the International Stroke Trial (IST)^[7]. In the CAST, 21,106 patients with acute ischemic stroke were enrolled in the clinical trial. Aspirin treatment (160 mg/d) was started within 48 h of the onset of a suspected acute ischemic stroke and continued in hospital for up to 4 wk. The results showed that aspirin treatment produced a significant reduction in mortality. Also, significantly fewer recurrent ischemic strokes oc-

curred in the aspirin group compared with the placebo-allocated group, but slightly more hemorrhagic strokes. In the IST, 19,435 patients with acute ischemic stroke within 48 h of symptom onset were enrolled. Anti-thrombotic therapy was started as soon as possible after stroke onset. The results showed that aspirin treatment significantly reduced death or recurrent ischemic strokes.

In addition to the two clinical trials, there are some completed clinical trials that investigated the antiplatelet strategy for acute ischemic stroke^[15,16]. In the FASTER (the fast assessment of stroke and transient ischemic attack to prevent early recurrence) trial, 392 patients with TIA or minor stroke within 24 h of symptom onset were enrolled^[15]. The trial was designed to investigate the effect of the combination of aspirin + clopidogrel versus aspirin monotherapy (with or without simvastatin) on the risk of recurrent stroke within 90 d after a TIA or minor stroke. The results showed the trend to lower recurrence or deterioration of stroke in the acute stage in the dual therapy group compared with mono antiplatelets. Another EARLY trial compared the safety and efficacy of aspirin + dipyridamole initiated within 24 h of stroke or TIA with that of aspirin + dipyridamole initiated after 7 d of aspirin monotherapy^[16]. The results showed that early initiation of aspirin plus extended-release dipyridamole did not lead to differing disability at 90 d compared with late initiation, but the proportion of patients in the early initiation group who had no or mild disability on day 90 was higher than that in the late initiation group. There was also no significant difference in the composite adverse event endpoint between the two groups. These studies demonstrated the efficacy and safety of combined antiplatelet treatment in acute ischemic stroke. Furthermore, the proposal also got support from the CLAIR study^[17]. The study aimed to investigate whether treatment with clopidogrel plus aspirin reduced the number of microembolic signals detected with transcranial doppler ultrasound compared with aspirin alone in patients with recent stroke. The results showed that combination therapy with clopidogrel and aspirin is more effective than aspirin alone in reducing microembolic signals in patients with predominantly intracranial symptomatic stenosis.

In other clinical trials aiming to investigate the secondary prevention for ischemic stroke, antiplatelet treatment was applied to some patients with acute ischemic stroke. For example, The ESPRIT Study compared aspirin with aspirin plus dipyridamole for secondary prevention after an ischemic stroke or transient ischemic attack^[18]. The study recruited 96 patients during the acute stage of ischemic stroke and TIA. The Prevention Regimen for Effectively Avoiding Second Strokes (PROFESS) trial is designed to compare extended-release dipyridamole plus aspirin *vs* clopidogrel in secondary stroke prevention^[19]. The study recruited 1360 patients during the acute stage of ischemic stroke. These studies also demonstrated the efficacy and safety of combined antiplatelet treatment. Furthermore, a recent meta-analysis study show that dual antiplatelet therapy appears to

be safe and effective in reducing stroke recurrence and combined vascular events in patients with acute ischemic stroke or transient ischemic attack compared with monotherapy^[20].

One powerful evidence comes from the Clopidogrel in High-risk Patients with Acute Nondisabling Cerebrovascular Events (CHANCE) study, which was just published in the *New England Journal of Medicine*^[21]. The study showed that among patients with a TIA or minor stroke who can be treated within 24 h after the onset of symptoms, the combination of clopidogrel and aspirin is superior to aspirin alone for reducing the risk of stroke in the first 90 d and does not increase the risk of hemorrhage.

BLEEDING EVENTS CAUSED BY ANTI-PLATELETS

When considering using more intensive antiplatelet therapy, the balance between reducing recurrent stroke and increasing major bleeding events needs to be considered. Previous studies found that combination antiplatelet increased the bleeding events compared with mono antiplatelets. For example, the PROfESS study found that aspirin plus extended-release dipyridamole produced more major hemorrhagic events than clopidogrel, including intracranial hemorrhage^[22]. The MATCH study showed that life-threatening bleeding was higher in the group receiving aspirin and clopidogrel *vs* clopidogrel alone^[23]. However, some studies found no difference between combination antiplatelets and mono antiplatelets, for example, the EARLY, FASTER and CLAIR trials^[15-17]. More importantly, the recent CHANCE study also did not find more bleeding events with combined antiplatelets compared with mono antiplatelets^[21]. The discrepancy could be due to the duration of antiplatelet treatment. Obviously, combination antiplatelets in the long-term trials such as PROfESS and MATCH^[22,23] produced more bleeding events. Thus, a time window of combination antiplatelets should exist at which risk outweighs benefit. Taken together, the evidence shows that combination antiplatelets in a short-term time window (maybe less than 3 wk) should be safe.

CONCLUSION

As discussed above, although IST and CAST provided the most elegant evidence for aspirin in the acute stage of ischemic stroke, growing evidence, including CHANCE, points to the superiority of combination antiplatelets over mono antiplatelets. On one hand, due to their different mechanisms of action, it is postulated that combined antiplatelet agents may provide different degrees of vascular protection. On the other hand, the superiority of combination therapy compared to monotherapy may be due to the synergy effects of different antiplatelet mechanisms in reducing vascular events. Thus, combined antiplatelet therapy, such as aspirin and clopidogrel, should be encouraged in the treatment of acute ischemic stroke.

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Atypical neurological symptoms associated with CGG expansions of the *FMR1* gene

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Abstract

More than 40 CGG expansions in the 5' noncoding region of the fragile X mental retardation 1 (*FMR1*) gene of the X chromosome give rise to several distinct clinical phenotypes, depending on the size of the expansion. First, more than 200 CGG expansions (full mutation) cause an inherited mental retardation called fragile X syndrome. Second, CGG expansions between 55 and 199 (premutation) cause a disorder called fragile X-associated tremor/ataxia syndrome (FXTAS) which typically includes intention tremor, ataxia and specific magnetic resonance imaging (MRI) findings. Indeed, it could develop parkinsonism although it usually shows features of postsynaptic parkinsonism. Finally, CGG expansions between 41 and 54 CGG (gray zone) are not considered normal but rarely develop abnormal neurological conditions. In this sense, the aim of this study is to report two atypical cases associated with CGG expansions of the *FMR1* gene. First, a *FMR1* premutation alleles carrier with an unusual phenotype, such as a presynaptic parkinsonism indistinguishable from Parkinson disease (PD) and a *FMR1* gray zone alleles carrier presented with neurological features, namely hand tremor, parkinsonism and ataxia, usually described in FXTAS, as well as orthostatic tremor. We conclude that,

on the one hand, *FMR1* premutation alleles might cause two phenotypes of parkinsonism, such as a presynaptic phenotype, indistinguishable from PD, and a postsynaptic phenotype, associated with clinical features of FXTAS. On the other hand, although *FMR1* gray zone alleles carriers were believed to have no abnormal neurological conditions, our study supports that they could develop FXTAS and other neurological disorders such as orthostatic tremor which has not been reported before associated with the *FMR1* gene.

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Key words: Fragile X-associated tremor/ataxia syndrome; Fragile X mental retardation; Gray zone; Parkinsonism; Orthostatic tremor

Core tip: In this study we report two atypical cases associated with CGG expansions of the fragile X mental retardation 1 (*FMR1*) gene. First, a *FMR1* premutation alleles carrier presented with a parkinsonism indistinguishable from Parkinson disease. Second, a *FMR1* gray zone alleles carrier presented with orthostatic tremor and neurological features associated with the fragile X-associated tremor/ataxia syndrome, such as hand tremor, parkinsonism and ataxia.

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INTRODUCTION

More than 40 CGG expansions in the 5' noncoding region of the fragile X mental retardation 1 (*FMR1*) gene

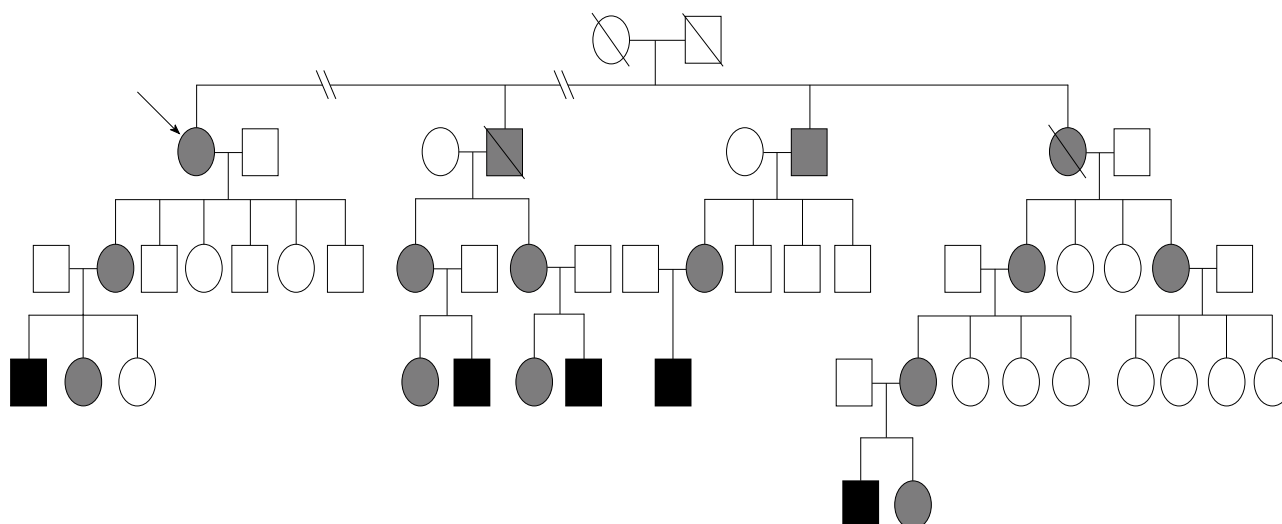


Figure 1 Pedigree structure of the patient 1 (arrow), they were 12 brothers and sisters, only those relevant are shown. Fragile X mental retardation (FMR1) CGG expanded alleles carriers are denoted as a grey filled symbol, they did not show any neurological disorder. Black filled symbols denote FMR1 CGG full mutation carriers affected by fragile X syndrome with mental retardation. Crossed symbol indicates deceased family members. Circle indicates females and square males.

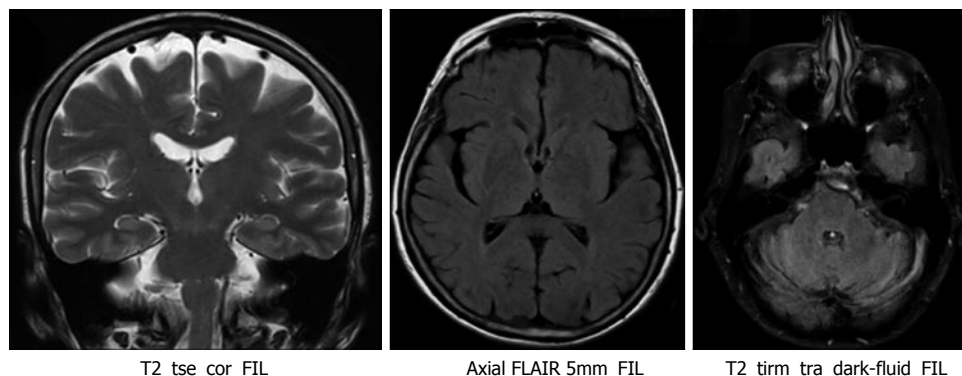


Figure 2 T2, TIR and FLAIR magnetic resonance imaging of the patient 1, it does not show abnormal findings. FLAIR: Fluid attenuated inversion recovery.

of the X chromosome give rise to several distinct clinical phenotypes, depending on the size of the expansion. First, fragile X syndrome is an inherited mental retardation caused by more than 200 CGG expansions in the *FMR1* gene (full mutation). Second, fragile X-associated tremor/ataxia syndrome (FXTAS) is a disorder described in premutation alleles carriers (55-199 CGG expansions) of the *FMR1* gene. Typically, it includes intention tremor, ataxia and specific magnetic resonance imaging (MRI) findings, although the spectrum of neurological disorders associated with the FXTAS is increasing^[1-3]. Finally, an interval of CGG expansions between 41 and 54 is known as gray zone which rarely develops abnormal neurological conditions. In this sense, the aim of this study is to report two atypical cases associated with CGG expansions of the *FMR1* gene. First, a FMR1 premutation alleles carrier with an unusual phenotype, such as a parkinsonism indistinguishable from Parkinson disease (PD) and a FMR1 gray zone alleles carrier presented with neurological features, namely hand tremor, parkinsonism, ataxia and orthostatic tremor.

CASE REPORT

Case 1

A 71-year-old woman presented with a 4-year history of progressive neurological disorder with tremor, bradykinesia and gait disorder. She was FMR1 alleles carrier (expansion of 82 CGG) with family history of fragile X syndrome (Figure 1). On examination, she had severe rest and postural tremor in her right hand and slight postural tremor in her left hand, severe bradykinesia and rigidity more marked on her right side, mild loss of postural stability and freezing. There was neither intention tremor nor ataxia. Indeed, patient did not show either autonomic dysfunction or Babinski sign or gaze palsy. Finally, She had significant improvement with 520 mg of levodopa (Movement Disorder Society-sponsored revision of the Unified Parkinson's Disease Rating Scale I 8 II 17 III ON 42 OFF 60 IV 0). These clinical findings fulfilled the United Kingdom PD Society Brain Bank clinical diagnostic criteria for idiopathic PD^[4].

Blood tests and cranial MRI were normal (Figure 2).

Table 1 Neuropsychological assessment of the patient 1

Test	Subtest	Percentile
Orientation		Normal
Attention and executive functions		
D2 test of attention	Processing elements	p45
	Omissions	p25
	Concentration	p40
Digit span (WMS-III)	Forward	p50
	Backward	p50
Verbal fluency	Semantic (animals)	p75
	Semantic (names)	p90
	Phonological (FAS)	p30
Trail making test	A	p50
	B	p20
Stroop		p50
Hanoi tower	3 disk	Normal
	4 disk	Not solved ¹
Similarities (WAIS)		p50
Wisconsin card sorting test	Categories completed	4 (P > 16)
	Failure to maintain set	p51
	Errors	p30
	Perseverative responses	p30
	Perseverative errors	p30
	Conceptual level responses	p20
BADS	Rule shift cards	p51
	DEX	Score 20
Episodic memory		
Logical memory I (WMS-III)	Recall unit (Story A + B)	p20
	Recall thematic (Story A + B)	p20
Logical memory II (WMS-III)	Recall unit (Story A + B)	p30
	Recall thematic (Story A + B)	p30
Word list I (WMS-III)	Recall total score	p50
	Short-delay recall	p85
Word list II (WMS-III)	Recall	p70
	Recognition	p85
Rey-osterrieth complex figure test	Recall	p50
Naming		
Boston naming test		p50
Praxis and visuo-constructive ability		
Rey-osterrieth complex figure test	Copy	p75
Gesture imitation		Normal
Complex sequencing		Normal

¹It shows slight deficits in speed information processing, in planning as well as in attention to designated targets and to inhibit responses. FAS: Foreign Agricultural Service; WAIS-III: Wechsler Adult Intelligence Scale-III; WAIS: Wechsler Adult Intelligence Scale; BADS: Behavioural Assessment of the Dysexecutive Syndrome.

DAT SCAN showed an asymmetric decrease of the striatum tracer uptake more marked on the left side (Figure 3). Neuropsychological assessment did not show cognitive decline. However, it showed slight subcortical deficits (Table 1).

Case 2

A 76-year-old man, FMR1 gray-zone alleles carrier (expansion of 51 CGG) presented with a 3-year history of hand tremor and severe gait disorder. On examination he had mild postural tremor in both hands, mild rigidity more marked in the left side, lost of postural stability, severe tremor in a standing position, ataxia and freezing.

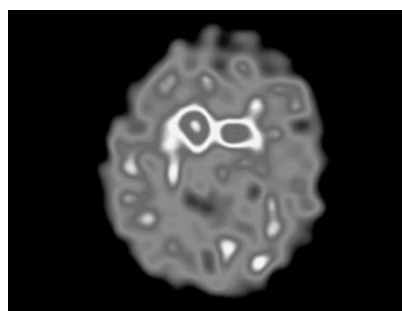


Figure 3 [¹²³I]FP-CIT DAT SCAN of the patient 1, it shows an asymmetric decrease of the striatum tracer uptake more marked on the left side.

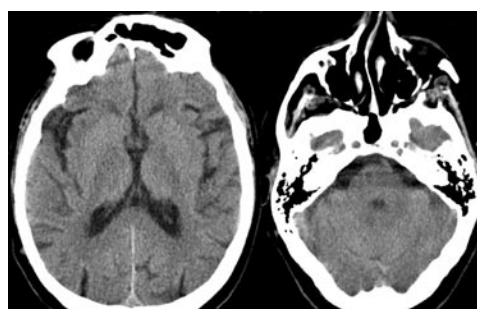


Figure 4 Cranial computed tomography of the patient 2, it does not show relevant findings.

He improved neither with 609 mg of levodopa nor with 1.5 mg of clonazepam. There were not cognitive decline.

Blood tests were normal. Cranial computed tomography and DAT SCAN were also normal (Figures 4 and 5). Electromyography corroborated high-frequency (14 Hz) tremor in a standing position in both legs which confirmed the diagnosis of orthostatic tremor.

DISCUSSION

Parkinsonism is often related to FXTAS, although most patients do not meet clinical criteria for PD^[1,2]. Indeed, in parkinsonism related to FXTAS, response to levodopa is often poor^[5,6], DAT SCAN is usually normal^[5] and Single Photon Emission Computed Tomography with iodo-benzamide has shown abnormal tracer uptake^[6]. It leads to conclude that parkinsonism in FXTAS is developed by dysfunction beyond the presynaptic nigrostriatal level^[5]. However, our first case, a parkinsonism with clinical features indistinguishable from PD, as similar cases previously reported^[2], without intention tremor or ataxia and with both good response to levodopa and abnormal DAT SCAN suggests damage restricted to the presynaptic level. Thus, although the relationship between these “presynaptic parkinsonisms” and the FMR1 premutation could be casual, it is necessary to consider that the FMR1 premutation might cause two phenotypes of parkinsonism, such as a presynaptic phenotype, indistinguishable from PD, and a postsynaptic phenotype, associated with clinical features of FXTAS. In fact, neuropathological features related to the *FMR1* gene without Lewy bodies

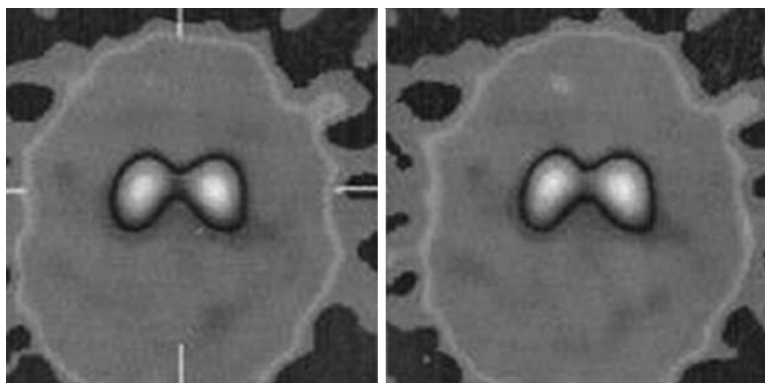


Figure 5 [^{123}I]FP-CIT DAT SCAN of the patient 2, it shows a symmetric and normal tracer uptake.

were found in a FMR1 premutation alleles carrier with clinical features of presynaptic parkinsonism^[7] which supported the relationship between that presynaptic parkinsonism and the *FMR1* gene as well as ruled out a concomitant PD. How to explain it remains unclear, it could reflect either different ranges of severity or different stages of the same process or the coexistence of the FMR1 premutation with others genes associated with PD^[8]. Thus, we suggest that FMR1 premutation should be considered in the differential diagnosis of PD, almost in those patients with family history of disorders related to the *FMR1* gene.

FMR1 gray zone alleles carriers were believed to have no abnormal neurological conditions. However, a recent report has suggested a relationship between FMR1 gray zone alleles carriers and FXTAS^[3]. In this sense, our second case, presented with usual clinical features of FXTAS, such as parkinsonism, hand tremor and ataxia, supports the fact that FMR1 gray zone alleles carriers could develop FXTAS. Indeed, orthostatic tremor has not been described before in association with the *FMR1* gene. Thus, the presence of orthostatic tremor in our second patient expands the clinical spectrum of neurological disorders associated with the *FMR1* gene. Future research should validate findings on a larger group of patients.

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- 3 **Tian D**, Araki H, Stahl E, Bergelson J, Kreitman M. Signature of balancing selection in Arabidopsis. *Proc Natl Acad Sci USA* 2006; In press

Organization as author

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Statistical data

Write as mean \pm SD or mean \pm SE.

Statistical expression

Express *t* test as *t* (in italics), *F* test as *F* (in italics), chi square test as χ^2 (in Greek), related coefficient as *r* (in italics), degree of freedom as *v* (in Greek), sample number as *n* (in italics), and probability as *P* (in italics).

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Use SI units. For example: body mass, *m* (B) = 78 kg; blood pressure, *p* (B) = 16.2/12.3 kPa; incubation time, *t* (incubation) = 96 h; blood glucose concentration, *c* (glucose) 6.4 ± 2.1 mmol/L; blood CEA mass concentration, *p* (CEA) = 8.6 $24.5 \mu\text{g/L}$; CO₂ volume fraction, 50 mL/L CO₂, not 5% CO₂; likewise for 40 g/L formaldehyde, not 10% formalin; and mass fraction, 8 ng/g, *etc.* Arabic numerals such as 23, 243, 641 should be read 23 243 641.

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Italics

Quantities: *t* time or temperature, *c* concentration, *A* area, *l* length, *m* mass, *V* volume.

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Biology: *H. pylori*, *E. coli*, *etc.*

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