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Pathologies of Calcium Channels

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Foreword

Calcium signaling is a highly versatile system responsible for regulating a large number of diverse cell functions in both excitable and non-excitable cells. It initiates a new life at the time of fertilization and then guides many of the subsequent developmental processes that orchestrate the emergence of all the multiple cell types that constitute all the organ systems of the human body. The enormous diversity and subtlety of these different signaling systems is quite remarkable. At synaptic endings, for example, a brief highly localized pulse of calcium can trigger transmitter release within microseconds, whereas global elevations of calcium lasting for seconds to minutes activate fertilization and metabolism in liver cells. To achieve this diversity, cells have access to an extensive calcium signaling toolkit from which they can select out and assemble unique signaling systems with the spatial and temporal properties necessary to control their particular cell-specific functions.

One of the features of these cell-type-specific calcium signaling systems is their relative stability. This stability depends on the turnover of signaling components being tightly regulated, which depends on a balance between the degradation of individual components and their replacement by ongoing transcription processes. Despite this stability, it is clear that these signaling systems are remarkably flexible and this has gradually come to light from knockout experiments where experimenters often are disappointed to find no apparent change in phenotype after removing an apparent key calcium signaling component. Such negative results are usually explained by invoking compensatory responses and are not investigated further. However, when the phenotype is examined in more detail, some surprising observations have emerged to indicate that when a particular component is removed, cells can interrogate their extensive toolkit to come up with an alternative closely related component that will restore the same function with varying degrees of effectiveness.

While this flexibility of the calcium signaling system enables organisms to survive in the face of changes in individual components, these compensatory changes are usually not perfect and the modified signaling systems often results in the emergence of a multiple diseased states. There is no better way of illustrating this point than to consider the contents of this remarkably complete anthology of calcium channel pathologies. It highlights how so many human diseases are caused by defects in calcium signaling systems in both excitable and non-excitable cells.

It is extraordinary to see that mutations in voltage-gated calcium channels have been linked to so many human pathologies and new examples are continuously appearing. For example, mutations in the *CACNA1C* gene that codes for the α_{1C} subunit of the $Ca_v1.2$ L-type calcium channel, which causes Timothy syndrome as described in this volume, has just been linked to a variety of psychiatric disorders such as schizophrenia and bipolar disorder. The insights gained from understanding the genetic basis of all these debilitating diseases are providing essential new information for our quest to understand how specific cell functions are controlled in complex organ systems such as the brain.

The editors, Norbert Weiss and Alexandra Koschak, have been extraordinary successful in bringing together so many of the major researchers in this rapidly expanding field. To have all this information in a single volume is a fantastic resource for both those workers actively involved in this exciting field and for those wishing to find out more about particular diseases.

Cambridge, UK   Sir Michael Berridge

Preface

Calcium ions (Ca^{2+}) represent one of the most versatile second messengers in eukaryotic cells, and changes in the intracellular Ca^{2+} concentration regulate many cellular physiological functions as diverse as gene transcription, muscle contraction, neurotransmission fertilization and more. In order to make use and regulate the amplitude, duration and subcellular localization of the Ca^{2+} signal, cells have developed a complex machinery, the Ca^{2+} signaling toolkit, initially introduced by Sir Michael Berridge, where Ca^{2+} channels play a key role in the initiation of the Ca^{2+} signal. Therefore, it is not surprising that mutations in Ca^{2+} channel genes, causing either a loss or a gain of channel function, can lead to serious (chronic) disorders. We have not only learned much within the last two decades regarding the structure/function basis of Ca^{2+} channels, but also the list of human diseases known to be associated with a defect in Ca^{2+} channels has grown considerably. This book aims to provide an informative and up-to-date account of our present understanding of human diseases associated to the various Ca^{2+} channels, from their molecular basis to the point of clinical pharmacology.

Voltage-Gated Ca $^{2+}$ Channels

Voltage-gated Ca^{2+} channels (VGCCs) are pore-forming plasma membrane proteins that convert an electrical signal into intracellular Ca^{2+} elevations. To date, 10 genes encoding the pore-forming subunits of mammalian VGCCs have been identified. 7 genes encode the high-voltage activated (HVA) channel subfamily (comprising L-type ($Ca_v1.1$ to $Ca_v1.4$), P/Q-type ($Ca_v2.1$), N-type ($Ca_v2.2$) and R-type ($Ca_v2.3$) channels) and 3 genes encode the low-voltage-activated (LVA) channel subfamily (composed exclusively of T-type ($Ca_v3.1$ to $Ca_v3.3$). In addition to the Ca_v pore-forming subunit, HVA channels contain auxiliary subunits: β (β_1 to β_4, a 55 kDa cytosolic protein of the membrane-associated guanylate kinase (MAGUK) family), $\alpha_2\delta$ ($\alpha_2\delta_1$ to $\alpha_2\delta_4$, a 170 kDa highly glycosylated extracellular protein with a single transmembrane domain), and in some cases γ (γ_1 to γ_8, a 33 kDa transmembrane protein). Calcium entry through neuronal $Ca_v2.1$ and $Ca_v2.2$ channels initiates fast neurotransmitter release at the central and peripheral synapses, respectively, by triggering the fusion of secretory vesicles

with the plasma membrane through actions on the SNARE synaptic protein complex. Therefore, small changes in their biophysical properties or subcellular expression result in alteration in neuronal excitability. In Chap. 1, Daniela Pietrobon discusses the implication of $Ca_v2.1$ channels in the pathophysiology of a rare form of migraine with aura, the familial hemiplegic migraine type I. Their implication in various forms of cerebellar ataxia in then discussed in Chap. 2 by Stephanie Schorge and Kinya Ishikawa. In contrast to Ca_v2 channels which are widely expression through the nervous system, $Ca_v1.3$ and $Ca_v1.4$ channels predominate in sensory cells of the inner ear and in retinal neurons, respectively, and have been linked to sensory disorders as discussed by Alexandra Koschak, Alexandra Pinggera, Klaus Schicker and Jörg Striessnig in Chap. 3. In contrast to HVA Ca^{2+} channels, LVA channels (referred as T-type channels or Ca_v3 channels), because of their particular activation threshold around the resting membrane potential of the cells, underlie neuronal burst-firing, often associated with spike-and-wave discharges observed on electroencephalography recordings during certain forms of epileptic seizures as discussed in Chap. 4 by Stuart M. Cain, Michael E. Hildebrand and Terrance P. Snutch. Finally, Annette Dolphin discusses the implication of the voltage-gated Ca^{2+} channel $\alpha_2\delta$-subunit in various neurological disorders and most importantly its clinical relevance as therapeutic target for gabapentinoid drugs (Chap. 5). In skeletal muscle, $Ca_v1.1$ functions as both a slowly-activating, voltage-gated L-type Ca^{2+} channel and as the voltage sensor that triggers the opening of ryanodine receptor Ca^{2+} release from the sarcoplasmic reticulum during excitation–contraction (EC) coupling (voltage-induced-Ca^{2+} release (VICR) mechanism). In Chap. 6, Amaury Francois Anne Pizzoccaro, Sophie Laffray and Emmanuel Bourinet review the current knowledge on the role of the LVA T-type channels in the perception and modulation of pain. In Chap. 7, Emma Matthews and Mike Hanna specifically discuss the implication $Ca_v1.1$ channels in hypokalemic periodic paralysis. Then, Viktor Yarotskyy and Robert T. Dirksen analyse its implication in malignant hyperthermia (Chap. 8), while Julien Fauré, Joël Lunardi, Nicole Monnier and Isabelle Marty investigate this functional coupling from the side of the ryanodine receptor type I (Chap. 9). In Chap. 10, Paul W. Wirtz, Maarten J. Titulaez and Jan J. G. M. Verschuuren discuss the molecular physiopathology of the Lambert-Eaton Mysathenic Syndrome, an autoimmune disease of the neuromuscular junction and the implication of $Ca_v2.1$ channels. In contrast to the skeletal $Ca_v1.1$ channel, cardiac $Ca_v1.2$ channels trigger the opening of the ryanodine receptor via a Ca^{2+}-induced-Ca^{2+}-release mechanism (CICR), and mutations in the gene encoding $Ca_v1.2$ channels have been linked to various cardiac syndrome, including the Timothy syndrome discussed by Hua Huang, Juejin Wang and Tuck Wah Soong in Chap. 11, and the Brugada syndrome discussed by Brett Simms in Chap. 12. In Chap. 13, Matteo Mangoni specifically discusses the implication of $Ca_v1.3$ channels in sinoatrial node dysfunction, and Stephan Herzig and Jan Matthes discuss in Chap. 14 how an alteration in the expression pattern of the VGCCs β-subunit and its consequence on the L-type current leads to an alteration of the cardiac function.

As previously mentioned, Ca^{2+} influx from VGCCs controls many cellular processes including cell growth and differentiation, and not surprisingly these channels have been implication in different form of cancer. This aspect is discussed in Chap. 15 by Maria Beatrice Morelli, Sonia Liberati, Consuelo Amantini, Matteo Santoni, Massimo Nabissi, Valerio Farfariello and Giorgio Santoni in the particular case of Ca_v3 channels. Finally, Ľubica Lacinová and Lucia Lichvárová close this section by providing an up-to-date picture of the clinically relevant drugs against VGCCs and associated diseases (Chap. 16).

Transient Receptor Potential Channels

In contrast to VGCCs, the Transient Receptor Potential (TRP) superfamily of cation channels displays greater diversity in activation mechanisms and selectivity. Based on the sequence homology, Mammalian TRP channels are divided into seven subfamilies: TRP Canonical (TRPC; TRPC1 to TRPC7), TRP Vanilloid (TRPV; TRPV1 to TRPV6), TRP Melastatin (TRPM; TRPM1 to TRPM8), TRP Polycystin (TRPP; TRPP2, TRPP3 and TRPP5), TRP Mucolipin (TRPML, TRPML1 to TRPML3), and TRP Ankyrin (TRPA; TRPA1). TRP channels are membrane spanning proteins that share the common features of six putative transmembrane segments and a non-selective permeability to cations. Although the cytoplasmic amino and carboxy termini are variable between TRP subfamily members, they contain different domains (i.e., ankyrin repeats, coiled–coiled, calmodulin binding sites and more) critical for channel regulation. Functional TRP channels form either homo- or heterotetramers assembled from identical or similar TRP subunits, and play critical roles in sensory physiology like vision, taste, olfaction, hearing, touch, but also thermo- and osmosensation. Although only few TRP members have been linked to hereditary diseases, the implication of TRP channels in the pathogenesis of many disease states has emerged. In Chap. 17, Shoichi Irie and Takahisa Furukawa present the physiological role of TRPM1 channel in retinal bipolar cells, and the association of human TRPM1 mutations with congenital stationary night blindness. In Chap. 18, Meredith C. Hermosura discusses how alteration in TRPM2 and TRPM7 functions in the immune system may lead to neurodegenerative diseases such as amyotrophic lateral sclerosis and Parkinsonism dementia. Christian Grimm and Math P. Cuajungco then highlight the cellular function of TRPML channels endolysosomal system and their implication in the lysosomal storage disorder mucolipidosis type IV (Chap. 19). As previously mentioned, activation of TRP channels present a great diversity in activation mechanisms, and are thus considered as polymodal cellular sensors. This diversity is particularly well illustrated by the implication of TRP channels in the signaling pathway of various painful stimuli, as discussed in Chap. 20 by Alexandre Denadai-Souza and Nicolas Cenac.

Besides their role in neurological disorders, the contribution of TRP channels has also been described in various muscular/skeletal and cardio-vascular disorders.

In Chap. 21, Jeremy M. Sullivan, Thomas E. Lloyd and Charlotte J. Sumner present the hereditary channelopathies (Charcot-Marie-Tooth disease, muscular atrophy, dysplasia and brachyolmia) caused by mutations in the gene encoding TRPV4 channel. Two chapters then discuss the role of TRPC channels in the cardiovascular system. In Chap. 22, Nadine Kirschmer, Kristina Lorenz and Petra Eder-Negrin explain how an alteration in TRPC channel activity may cause cardiac hypertrophy, whereas in Chap. 23, Michael Poteser, Sarah Krenn, and Klaus Groschner report on the contribution of these channels in the remodeling of the vascular system.

The broad implication of TRP channels in multiple diseased states in then further illustrated with the role of TRPP2 channel in hereditary polycystic kidney disease in Chap. 24 by Andrew Streets and Albert Ong. In Chap. 25, Daniel Landau and Hanna Shalev further illustrate the role of TRP channels in kidney related diseases and discuss the contribution of TPM6 channel in the development of hypomagnesemia and hypocalcemia.

Cell proliferation and differentiation highly relies on the Ca^{2+} signal, and alteration of the Ca^{2+} homeostasis often leads to disruption of normal cell cycle and, as a result, onset and progression of cancer. This aspect is further discussed in the context of prostate cancer by George Shapovalov, Roman Skryma and Natalia Prevarskaya in Chap. 26.

Finally, Ulrich Wissenbach concludes this section by discussing the pharmacological relevance of TRP channels as therapeutic targets (Chap. 27).

Ligand-Gated Channels

The third part of this book presents the channelopathies related to ligand-gated channels. As VGCCs and TRP Channels, Ligand-Gated Ca^{2+} channels form a group of transmembrane proteins, which open in response to the binding of a chemical messenger (ligand), allowing ions to pass through the membrane. This part presents the few Ca^{2+} channelopathies caused by alteration of the activity of ligand-gated Ca^{2+} channels.

IP3 Receptors

Inositol 1,4,5-trisphosphate (IP3) receptors (IP3Rs) are localized to intracellular membranes (such as the endoplasmic reticulum, the mitochondria and the nucleus) where they mediate the mobilization of intracellular Ca^{2+} stores following binding of IP3. There are three IP3 receptors subtypes (IP3R1, IP3R2 and IP3R3) exist as homo- and heterotetramers, and play an important role in intracellular Ca^{2+} signaling in a variety of cell types including nerve cells. In Chap. 28, Masayoshi Tada, Masatoyo Nishizawa and Osamu Onodera discuss the implication of these

receptors in the development of neurodegenerative disorders including spinocerebellar ataxias, and Huntington's and Alzheimer's diseases.

Ryanodine Receptors

Ryanodine Receptors (RyRs), localized in the membrane of the endoplasmic reticulum and as IP3Rs they contribute to the release of Ca^{2+}. There exist three RyR isoforms (RyR1 to RyR3). Whereas RyR1 is primarily expressed in skeletal muscle and shows a voltage-depend activation mode (see Chap. 9), RyR2 and RyR3, are predominantly expressed in the heart and the brain, respectively, and are activated by a local rise in cytosolic Ca^{2+} thereby contributing to the CICR. In the heart, Ca^{2+} release via RyR2 plays a critical role in muscle contraction, and mutations in *RyR2* have been linked to cardiac disorders. In Chap. 29, Ineke Nederend, Christian van der Werf and Arthur A. M. Wilde discuss the molecular mechanisms by which RyR2 channel contribute to the development of an inherited arrhythmia syndrome, the catecholaminergic polymorphic ventricular tachycardia.

P2X Receptors

Purinergic (P2X) receptors are plasma membrane proteins that open in response to changes in extracellular ATP. Each receptor is made up of a trimer of identical subunits (P2X1 to P2X7), or can also exist as heteromer (with the exception of P2X6 which cannot form functional homomeric receptor). They share the common structure of two transmembrane domains, a large extracellular loop and intracellular amino- and carboxy-termini. P2X receptors are widely expressed throughout the body, including the central and peripheral nervous system, smooth muscle, heart and leukocytes, and are involved in a range of physiological processes such as modulation of synaptic transmission, vascular tone, cardiac rhythm and contractility, but also nociception as described in Chap. 30 by François Rassendren and Lauriane Ulmann. In Chap. 31, Melissa L. Barron, Eryn L. Werry, Iain S. McGregor and Michael Kassiou specifically elucidate the role of P2X7 receptor in depressive and bipolar disorders.

NMDA Receptors

N-methyl-*D*-aspartate (NMDA) receptors are members of the ionotropic class of glutamate receptors. They consist of two NR1 subunits combined with two NR2 (NR2A to NR2D) or NR3 (NR3A or NR3B) subunits. Activation of NMDA receptors requires binding of glutamate or aspartate, as well as binding of co-

agonist glycine for the efficient opening of the channel. NMDA receptors are non-selective cation permeable channels, including Na^+, Ca^{2+} and K^+. Calcium influx through NMDA receptors is believed to be critical in synaptic plasticity, and it is not surprising that his involvement in the pathophysiology of epilepsy has been proposed, as discussed by Mehdi Ghasemi and Ahmad Reza Dehpour in Chap. 32.

Nicotinic Acetycholine Receptors

Nicotinic acetycholine receptors (nAChRs) are plasma membrane cholinergic receptors, expressed in many neurons and on the postsynaptic side of the neuro-muscular junction. Each receptor is composed of pentameric combinations of subunits (α1 to α10, β1 to β4, δ, ε and γ), and opening of the channel is triggered by the binding of the neurotransmitter acetycholine (Ach). Neuronal nicotinic receptors are found in the central nervous system and in autonomic ganglia where they regulate neurotransmitter release, cell excitability and neuronal integration. In contrast, nicotinic receptors located at the neuromuscular junctions of somatic muscles are responsible for muscular contraction. The implication of nAChRs in muscle pathophysiology is illustrated in Chap. 33 by Francesca Grassi and Sergio Fucile in the context of congenital myasthenic syndromes.

CatSper Channels

CatSper channels are sperm-specific cations channels. There are four CatSper isoforms (CatSper1 to CatSper4) surrounded by at least three auxiliary subunits (α, β and γ), and consist of six putative transmembrane segments with intracellular amino- and carboxy-termini. These channels are weakly voltage-dependent and are activated by intracellular alkalinization and several extracellular ligands, i.e., progesterone and prostaglandin E in human spermatozoa. CatSper channels are essential for hyperactivation of sperm motility required for fertility. Hence, in their chapter, Takuya Nishigaki, Ana Laura González Cota and Gerardo José Orta Salazar present the functional importance of CatSper channels in sperm physiology and discuss their implication in male infertility (Chap. 34).

Together, these expert contributions provide the current state-of-play in this exciting and medically important area of physiology.

Calgary, Canada   Norbert Weiss
Vienna, Austria   Alexandra Koschak

Acknowledgments

First, we would like to warmly thank the authors who agreed to contribute to this book, especially in these *impact factor* dominated days. We feel happy to see that the motivational factor for commitment and willingness to contribute a chapter has been the sole and unique scientific interest of bringing together the current state-of-play in this exciting area of calcium channelopathies in a single volume. Writing a book chapter is not an easy task and we apologize to the authors for stressing them by sending repeated reminders. This was the "price to pay" to publish an up-to-date volume in this rapidly expanding field.

We also thank the referees who generously agreed to review the chapters, and have helped the authors in this difficult task of writing a book chapter and, who have as well as contributed to the project giving their valuable suggestions.

At the time of writing this, we realize that more than a year has passed since we started to discuss this book project. For us it has been a rewarding scientific and human experience and we believe that all the readers, whether basic researchers, clinicians, graduate students, or hopefully also general readers interested by this exciting and medically important area of physiology will find something valuable in this book.

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Norbert Weiss
Alexandra Koschak
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Contents

Part I
Pathologies of Voltage-Gated Calcium Channels

Chapter 1
Ca$_v$2.1 Channels and Migraine

Daniela Pietrobon

Abstract Migraine is a common disabling brain disorder whose key manifestations are recurrent attacks of unilateral headache that may be preceded by transient neurological aura symptoms. Missense mutations in CACNA1A, the gene that encodes the pore-forming α_1 subunit of human voltage-gated Ca$_v$2.1 (P/Q-type) calcium channels, cause a rare form of migraine with aura (familial hemiplegic migraine type 1: FHM1). This chapter, first, briefly summarizes current understanding of the pathophysiological mechanisms that underlie migraine headache, migraine aura, and the onset of a migraine attack. Then, the chapter describes and discusses (i) the functional consequences of FHM1 mutations on the biophysical properties of recombinant human Ca$_v$2.1 channels and native Ca$_v$2.1 channels in neurons of knockin mouse models carrying the mild R192Q or severe S218L mutations in the orthologous gene, and (ii) the functional consequences of these mutations on neurophysiological processes in the cerebral cortex and trigeminovascular system thought to be involved in the pathophysiology of migraine (including cortical spreading depression, cortical synaptic transmission, and several trigeminal ganglion neuron functions) and the insights into migraine mechanisms obtained from the alterations of these processes in FHM1 knockin mice.

1.1 Introduction

Migraine is an episodic neurological disorder characterized by recurrent attacks of typically throbbing and unilateral, often severe, headache with certain associated features such as nausea, phonophobia, and/or photophobia; in a third of patients the headache is preceded by transient neurological symptoms, that are most

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frequently visual, but may involve other senses (migraine with aura: MA) (Lipton et al. 2004). Migraine is remarkably common (e.g., it affects 17 % of females and 8 % of males in the European population (Stovner and Hagen 2006), very costly (EUR 1.8 billion per year in Europe) (Olesen et al. 2012), and disabling (one of the 20 most disabling diseases according to the World Health Organization: (Leonardi et al. 2005).

Migraine is a complex genetic disorder, with heritability estimates as high as 50 % and a likely polygenic multifactorial inheritance (de Vries et al. 2009). Although recent genome-wide association studies have identified a few risk factors for migraine (Anttila et al. 2010; Chasman et al. 2011; Freilinger et al. 2012), most of our current molecular understanding comes from studies of familial hemiplegic migraine (FHM), a rare monogenic autosomal dominant form of MA (Pietrobon 2007; de Vries et al. 2009; Russell and Ducros 2011).

Three FHM causative genes have been identified, all encoding ion channels or transporters (Ophoff et al. 1996; De Fusco et al. 2003; Dichgans et al. 2005). FHM type 1 (FHM1) is caused by missense mutations in CACNA1A (chromosome 19p13), that encodes the pore-forming α_1 subunit of human voltage-gated Ca$_v$2.1 (P/Q-type) calcium channels (Ophoff et al. 1996).

The FHM1 mutations produce substitutions of conserved amino acids in important functional regions of the Ca$_v$2.1 channel including the pore lining and the voltage sensors (see (Pietrobon 2010; Russell and Ducros 2011; Pietrobon 2013) for recent reviews and references). Figure 1.1 shows the location of FHM1 (and a few sporadic hemiplegic migraine, SHM1) mutations in the secondary structure of the Ca$_v$2.1α_1 subunit.
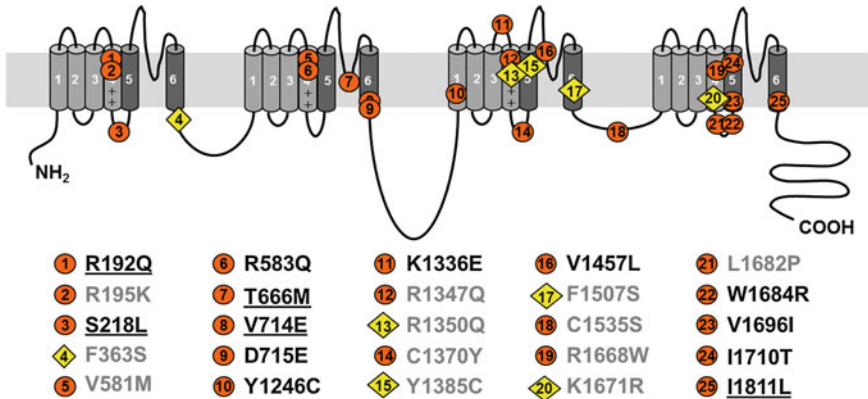
Fig. 1.1 Location of FHM1 mutations in the secondary structure of the Ca$_v$2.1α_1 subunit. FHM1 mutations are indicated with *circles*; a few SHM1 mutations not found in either controls or parents and affecting conserved amino acids are indicated with rhombs. Reference sequence Genbank Ac. N. X99897. The mutations written in *black* are those whose effects on the biophysical properties of recombinant Ca$_v$2.1 channels have been studied in heterologous expression systems; the mutations underlined are those whose effects have been studied also in transfected neurons from Ca$_v$2.1$^{-/-}$ mice expressing human Ca$_v$2.1α_1 subunits

FHM1 can be considered a model for the common forms of migraine because, apart from the motor weakness or hemiparesis during aura (and the possibly longer aura duration), typical FHM attacks resemble MA attacks (Lipton et al. 2004) and both types of attacks may alternate in patients and co-occur within families (Pietrobon 2007; de Vries et al. 2009; Russell and Ducros 2011). However, several FHM1 families show permanent cerebellar symptoms (such as slowly progressive cerebellar ataxia and/or nystagmus with cerebellar atrophy in some cases), that are usually not observed in common migraines. Pure FHM1 and FHM1 with cerebellar symptoms are usually associated with different mutations. Moreover, in addition to typical attacks, patients of some FHM1 families can have atypical severe attacks with signs of diffuse encephalopathy, impairment of consciousness (coma) or confusion, fever, prolonged hemiplegia lasting several days, and in a few cases seizures (Ducros et al. 2001; Russell and Ducros 2011).

Two different FHM1 knockin mouse models have been generated by introducing the human R192Q and S218L mutations into the orthologous *cacna1a* Ca$_v$2.1 channel gene (van den Maagdenberg et al. 2004, 2010). Whereas mutation R192Q causes in humans pure FHM, mutation S218L causes a particularly dramatic clinical syndrome, that may consists of, in addition to attacks of hemiplegic migraine, slowly progressive cerebellar ataxia and atrophy, epileptic seizures, coma or profound stupor and severe, sometimes fatal, cerebral oedema which can be triggered by only a trivial head trauma (Kors et al. 2001; van den Maagdenberg et al. 2010). Whereas homozygous R192Q and heterozygous S218L knockin mice did not exhibit an overt phenotype, homozygous S218L mice exhibited mild permanent cerebellar ataxia, spontaneous attacks of hemiparesis and/or (sometimes fatal) generalized seizures, and brain edema after only a mild head impact, thus modeling the main features of the severe S218L clinical syndrome (van den Maagdenberg et al. 2004, 2010).

Ca$_v$2.1 channels are located in presynaptic terminals and somatodendritic membranes throughout the mammalian brain and spinal cord (Westenbroek et al. 1995), and play a prominent role in initiating action potential (AP)-evoked neurotransmitter release at central nervous system synapses (Pietrobon 2005a). At many central synapses P/Q-, N-, and R-type Ca^{2+} (Ca) channels cooperate in controlling neurotransmitter release, but P/Q-type channels have a dominant role, partly because of a more efficient coupling to the exocytotic machinery (Mintz et al. 1995; Wu et al. 1999; Li et al. 2007). Moreover, at many central synapses, there is a developmental change in the Ca channel types mediating synaptic transmission, whereby the relative contribution of P/Q-type channels to release increases with postnatal age, until release becomes exclusively dependent on P/Q-type channels (Iwasaki et al. 2000). Among the presynaptic Ca channels, Ca$_v$2.1 channels are unique also in their capacity of interacting with and being modulated in a complex manner by a number of Ca-binding proteins ((Catterall and Few 2008) for review and references). As a result, Ca$_v$2.1 channels may exhibit both Ca-dependent inactivation and Ca-dependent facilitation. Ca-dependent regulation of presynaptic Ca$_v$2.1 channels may play a crucial role in short-term synaptic plasticity during trains of action potentials (Inchauspe et al. 2004; Mochida et al. 2008; Lin et al. 2012).

The somatodendritic localization of $Ca_v2.1$ channels points to additional postsynaptic roles, e.g., in neural excitability (Pineda et al. 1998; Womack et al. 2004), gene expression (Sutton et al. 1999), and cell survival (Fletcher et al. 2001).

$Ca_v2.1$ channels are expressed in all brain regions that have been implicated in the pathogenesis of migraine and/or migraine pain (cf. next Section), including the cerebral cortex (see Sect. 1.5.1), the trigeminal ganglia, and brainstem nuclei involved in the central control of nociception (see Sect. 1.5.2). Their expression is particularly high in the cerebellum, a feature that explains the cerebellar symptoms caused by several FHM1 mutations ((Pietrobon 2010) and Chapter 2 in this book).

1.2 Migraine Pathophysiology

It is generally recognized that most migraine attacks start in the brain, as suggested by the premonitory symptoms (such as difficulty with speech and reading, increased emotionality, sensory hypersensitivity), that in many patients are highly predictive of the attack although occurring up to 12 h before it, and as suggested also by the nature of some typical migraine triggers such as stress, sleep deprivation, oversleeping, hunger, prolonged sensory stimulation; moreover, psychophysical and neurophysiological studies have provided clear evidence that in the period between attacks migraineurs show hypersensitivity to sensory stimuli and abnormal processing of sensory information (see (Pietrobon and Striessnig 2003; Aurora and Wilkinson 2007; Coppola et al. 2007; Vecchia and Pietrobon 2012; Pietrobon and Moskowitz 2013) for reviews and references).

The neurophysiologial correlate of migraine aura is cortical spreading depression (CSD), a self-sustaining slowly propagating (2–5 mm/min) wave of nearly complete depolarization of a sizable population of brain cells that lasts 1–2 minutes (see (Lauritzen 1994; Pietrobon and Striessnig 2003; Ayata 2009; Charles and Brennan 2009) for reviews and references). CSD can be induced in healthy brain tissue by stimulation of the cerebral cortex with intense depolarizing stimuli. Although the mechanisms of initiation and propagation of CSD remain unclear, the initiation of the positive feedback cycle that ignites CSD and almost zeroes the neuronal membrane potential is thought to depend on the local increase of the extracellular concentration of K^+ ions above a critical value and on the activation of a net inward current at the pyramidal cell dendrites (Somjen 2001).

It is generally believed that the headache phase of migraine begins with the activation and sensitization of trigeminal sensory afferents that innervate cranial tissues, in particular the meninges and their large blood vessels; this then leads to sequential activation (and, in most patients, sensitization) of second and third order trigeminovascular neurons (in the trigeminal nucleus caudalis and specific thalamic nuclei, respectively), which in turn activate different areas of the brainstem and forebrain resulting in pain and other symptoms of migraine; the bidirectional signaling between many of these areas contributes to the complexity of the

symptoms (see (Olesen et al. 2009; Levy 2010; Akerman et al. 2011; Pietrobon and Moskowitz 2013) for recent reviews and references).

Most dural trigeminovascular afferents can be activated and sensitized by an inflammatory soup and are capsaicin-sensitive, thus exhibiting properties characteristic of nociceptors in other tissues (Strassman et al. 1996; Bove and Moskowitz 1997; Strassman and Levy 2006; Levy et al. 2007; Levy 2010; Vaughn and Gold 2010; Fioretti et al. 2011; Yan et al. 2011). Activation of meningeal nociceptors in vivo leads to release of vasoactive proinflammatory peptides such as CGRP and substance P from their peripheral nerve endings, that produce vasodilation of meningeal blood vessels (mainly due to CGRP), plasma extravasation and local activation of dural mast cells, with ensuing release of cytokines and other inflammatory mediators (neurogenic inflammation) (Waeber and Moskowitz 2005; Levy 2010). Measurements of CGRP levels into the external and internal jugular venous blood have provided evidence that CGRP is released during migraine attacks, and several findings support a pivotal role of CGRP in migraine, including the efficacy of CGRP receptor antagonists in migraine treatment (reviewed in (Recober and Russo 2009; Villalon and Olesen 2009; Ho et al. 2010)). However, the mechanism of actions of CGRP during a migraine attack and the exact sites of action of CGRP receptor antagonists remain unclear and controversial. The localization of CGRP receptors in the trigeminovascular system points to multiple possible mechanisms at both peripheral and central sites (Lennerz et al. 2008).

Although a large body of indirect evidence supports the idea that a sterile meningeal inflammation is a key mechanism that may activate and sensitize perivascular meningeal afferents and lead to migraine pain (reviewed in (Waeber and Moskowitz 2005; Levy 2009, 2010; Vecchia and Pietrobon 2012; Pietrobon and Moskowitz 2013)), the endogenous processes that promote meningeal inflammation and the mechanisms underlying the sustained activation and sensitization of the trigeminovascular system resulting in the typical long-lasting throbbing headache remain unclear.

The mechanisms of the primary brain dysfunction(s) leading to the onset of a migraine attack, to CSD susceptibility and to episodic activation of the trigeminovascular pain pathway remain largely unknown and remain the major open issues in the neurobiology of migraine.

Increasing evidence from animal studies support the idea that CSD can activate trigeminal nociception and thus trigger the headache mechanisms (Vecchia and Pietrobon 2012; Pietrobon and Moskowitz 2013). A direct nociceptive effect of CSD has been demonstrated by the finding that a single CSD can lead to a long-lasting increase in ongoing activity of dural nociceptors and central second order trigeminovascular neurons (Zhang et al. 2010, 2011), and evokes alterations in the meninges and brainstem consistent with the development of pain (Bolay et al. 2002). An interesting recent study provided evidence that CSD leads to opening of neuronal pannexin channels and consequent release of proinflammatory mediators, which initiates a parenchymal inflammatory response leading to sustained release of inflammatory mediators from glia limitans; this CSD-induced inflammatory

cascade may provide the sustained stimulus required for sensitization of trigeminal nociceptors and lasting pain (Karatas et al. 2013).

The idea that CSD is noxious and may trigger headache is indirectly supported by the finding that the electrical stimulation threshold for induction of CSD in the rat cortex increases after chronic treatment with five different migraine prophylactic drugs, that are effective in reducing the frequency of migraine attacks (both with and without aura) (Ayata et al. 2006). Moreover, further support to the view of CSD as a key migraine trigger has been provided by the analysis of experimental CSD in FHM knockin mouse models, that revealed a lower electrical stimulation threshold for CSD induction and a higher velocity of CSD propagation in both FHM1 and FHM2 knockin mice (van den Maagdenberg et al. 2004, 2010; Eikermann-Haerter et al. 2009b; Leo et al. 2011) (see Sect. 1.5.1). However, despite the strong support provided by animal studies, the idea that CSD may initiate the headache mechanisms in migraine is not generally accepted, mainly because it seems unable to explain some clinical observations, in particular the lack of a fixed relationship between aura and headache (reviewed in (Pietrobon and Striessnig 2003; Charles 2010; Vecchia and Pietrobon 2012; Pietrobon and Moskowitz 2013)).

The analysis of interictal cortical excitability using psychophysical, electrophysiological, and brain imaging methods has produced contradictory findings and interpretations regarding the mechanisms underlying the abnormal processing of sensory information in migraineurs (see (Pietrobon and Striessnig 2003; Aurora and Wilkinson 2007; Coppola et al. 2007; Vecchia and Pietrobon 2012) for reviews and references). Interestingly, recent transcranial magnetic stimulation studies in MA patients point to deficient regulatory mechanisms of cortical excitability and consequent reduced ability to dynamically maintain the cortical excitatory-inhibitory balance and to prevent excessive increases in cortical excitation, rather than merely hypo- or hyperexcitability, as the mechanisms underlying abnormal sensory processing (Antal et al. 2008; Conte et al. 2010; Siniatchkin et al. 2012). The molecular and cellular mechanisms underlying the abnormal regulation of cortical function and the increased susceptibility to CSD in migraine remain largely unknown. As discussed in Sect. 1.4, the functional analysis of FHM knockin mouse models supports the view of migraine as a disorder of brain excitability characterized by deficient regulation of the cortical excitatory-inhibitory balance, and gives insights into the possible underlying molecular and cellular mechanisms and their relationship to CSD susceptibility.

1.3 Effect of FHM1 Mutations on the Biophysical Properties of $Ca_v2.1$ Channels

The analysis of the P/Q-type calcium current in different neurons (including cortical and TG neurons) of R192Q and S218L FHM1 knockin mice revealed gain-of-function of the $Ca_v2.1$ current in a wide range of relatively mild depolarizations,

reflecting shifted activation of mutant Ca$_v$2.1 channels to more negative voltages (van den Maagdenberg et al. 2004, 2010; Tottene et al. 2009; Inchauspe et al. 2010; Fioretti et al. 2011 Gao et al. 2012). P/Q current densities were similar in knockin and wild-type (WT) neurons at higher voltages (that elicit maximal Ca$_v$2.1 channel open probability), indicating similar densities of functional Ca$_v$2.1 channels (van den Maagdenberg et al. 2004, 2010; Tottene et al. 2009; Fioretti et al. 2011; Gao et al. 2012). The shift to lower voltages of Ca$_v$2.1 channel activation and the gain-of-function of the neuronal Ca$_v$2.1 current were about twice as large in homozygous compared to heterozygous knockin mice, revealing an allele-dosage effect consistent with dominance of the mutation in FHM1 patients (van den Maagdenberg et al. 2010). In correlation with the severity of the clinical phenotype, the gain-of-function of the P/Q Ca current at low voltages was larger in cerebellar granule cells of S218L than R192Q knockin mice (van den Maagdenberg et al. 2004, 2010). Two photon microscopy in cerebellar slices of S218L knockin mice revealed increased AP-evoked Ca transients in individual synaptic terminals of cerebellar granule cells, a finding consistent with increased open probability of mutant pre-synaptic Ca$_v$2.1 channels (Adams et al. 2010), that may be due to shifted activation to lower voltages, as shown for the somatic channels (van den Maagdenberg et al. 2010), and/or to basal Ca-dependent facilitation of mutant channels as suggested by the authors (Adams et al. 2010) (see sect. 1.4).

The gain-of-function effect of the FHM1 mutations on native neuronal mouse Ca$_v$2.1 channels is in agreement with the increased open probability (mainly due to shifted activation to lower voltages) of recombinant human Ca$_v$2.1 channels carrying eight different FHM1 mutations (including R192Q and S218L), that was revealed by single channel recordings (Hans et al. 1999; Tottene et al. 2002; Tottene et al. 2005; Catterall et al. 2008) (our unpublished observations). The larger gain-of-function of the P/Q Ca current in neurons of S218L compared to R192Q knockin mice at low voltages is in agreement with the lower threshold of activation of human S218L Ca$_v$2.1 compared to human R192Q Ca$_v$2.1 channels (Tottene et al. 2005).

Consistent (and significant for the majority of the 13 FHM1 mutations analyzed so far, shown in black in Fig. 1.1) shifts to lower voltages of activation of recombinant mutant channels were also revealed by measurements of whole-cell current in heterologous expression systems (Kraus et al. 1998, 2000; Hans et al. 1999; Tottene et al. 2002, 2005; Melliti et al. 2003; Mullner et al. 2004; Weiss et al. 2008; Adams et al. 2009; Serra et al. 2009; Garza-Lopez et al. 2012) and transfected neurons (where five FHM1 mutations, underlined in Fig. 1.1, were analyzed) (Tottene et al. 2002, 2005). Moreover, these studies have shown that the FHM1 mutations may also alter the inactivation properties of Ca$_v$2.1 channels; the effects were different depending on the mutation and, e.g., resulted in increased, decreased, or unaltered inactivation during train of pulses, depending on the mutation.

For certain mutations, the specific alterations of channel gating, including the magnitude (or even the presence) of the shift to lower voltages of channel activation, depended on the Ca$_v$2.1α_1 splice variant and the type of coexpressed β subunit (Mullner et al. 2004; Adams et al. 2009). The use of different Ca$_v$2.1 splice variants and/or β subunits might explain some discrepancies existing in the

literature regarding the functional effects of some FHM1 mutations in transfected cells: e.g., the T666 M mutation was shown to produce a negative shift of $Ca_v2.1$ activation in (Kraus et al. 1998; Tottene et al. 2002) in contrast, a lack of effect on activation gating was reported in (Barrett et al. 2005; Tao et al. 2012).

Moreover, the expression of specific $Ca_v2.1$ splice variants and/or auxiliary subunits in different neurons might underlie the recent finding of a differential effect of the R192Q mutation on activation gating of native P/Q-type Ca channels in different TG neurons of knockin mice (Fioretti et al. 2011). In small capsaicin-insensitive TG neurons characterized by expression of T-type Ca currents (CI-T neurons) there was a larger P/Q-type Ca current density following mild depolarizations and a larger AP-evoked P/Q calcium current when compared to CI-T neurons from WT mice. In striking contrast, the P/Q-type current density, voltage dependence, and kinetics were not altered by the FHM1 mutation in small capsaicin-sensitive neurons of R192Q knockin mice (Fioretti et al. 2011).

Interestingly, while in cortical pyramidal cells of R192Q KI mice the shift to lower voltages of $Ca_v2.1$ channel activation resulted in increased AP-evoked Ca current, a similar shift of mutant R192Q channels at the Calyx of Held synaptic terminals did not alter the AP-evoked Ca current (Inchauspe et al. 2010). The different durations of the AP in pyramidal cells and Calyx (1.8 vs 0.44 ms AP half width) may largely explain the differential effects of the FHM1 mutation on the AP-evoked Ca current; in fact, the AP-evoked Ca current became larger in knockin compared to WT Calyx terminals when the longer duration pyramidal cell AP was used as depolarizing stimulus (Inchauspe et al. 2010).

The findings of (Fioretti et al. 2011) and (Inchauspe et al. 2010) have important general implications for familial migraine mechanisms, in that neuron subtype-specific (and/or subcellular compartment-specific) alterations of $Ca_v2.1$ channels and/or AP-evoked Ca influx may help to explain why a mutation in a Ca channel that is widely expressed in the nervous system (Westenbroek et al. 1995) produces the specific neuronal dysfunctions leading to migraine (see Sect. 1.5).

In contrast with the unaltered densities of functional $Ca_v2.1$ channels in neurons of FHM1 knockin mice (van den Maagdenberg et al. 2004, 2010; Tottene et al. 2009; Fioretti et al. 2011), overexpression of recombinant mutant $Ca_v2.1$ channels in transfected cells leads to alterated (decreased for most mutations) density of functional channels in the membrane, that results in reduced maximal whole-cell P/Q-type Ca current in cells expressing mutant channels (Hans et al. 1999; Tottene et al. 2002; Cao et al. 2004; Barrett et al. 2005; Cao and Tsien 2005; Tottene et al. 2005; Tao et al. 2012), including neurons expressing R192Q and S218L $Ca_v2.1$ channels (Tottene et al. 2002; Cao et al. 2004; Cao and Tsien 2005; Tottene et al. 2005) (cf. Fig. 3 in (Pietrobon 2010)).

1.4 Effect of FHM1 Mutations on Modulation of Ca$_v$2.1 Channels

Studies of recombinant human Ca$_v$2.1 channels carrying five different FHM1 mutations have shown that the migraine mutations reduce G-protein-mediated inhibitory modulation of Ca$_v$2.1 channels (Melliti et al. 2003; Weiss et al. 2008; Serra et al. 2009; Garza-Lopez et al. 2012), an effect that may lead to further gain-of-function of Ca influx through mutant channels during neuromodulation. The only study that investigated voltage-dependent G-protein modulation using physiological-like stimuli revealed a reduced G$\beta\gamma$-mediated inhibition of the Ca current evoked by AP-like stimuli and a reduced facilitation of the Ca current carried by mutant channels after a train of short depolarizing pulses in cells overexpressing G$\beta\gamma$ subunits; the reduction of G-protein modulation was dependent on the duration of the AP, being larger with APs of longer duration (Serra et al. 2009).

Indirect evidence of reduced G-protein-mediated inhibition of native neuronal Ca$_v$2.1 channels in R192Q knockin mice has been recently obtained in a study investigating the effect of the FHM1 mutation on sleep regulation by adenosine receptors in vivo; in fact, the FHM1 mouse models showed reduced responsiveness to either inhibition or enhancement of adenosine receptor activation (Deboer et al. 2013).

Besides G-protein modulation, the FHM1 mutations may also reduce Ca-dependent facilitation (CDF) of Ca$_v$2.1 channels. CDF of recombinant Ca$_v$2.1 channels was largely reduced by the R192Q mutation and almost eliminated by the S218L mutation (Adams et al. 2010), suggesting that the increase in the open probability of mutant channels (larger for S218L than R192Q: (Tottene et al. 2005)) partially or completely occludes further facilitation. CDF of native Ca$_v$2.1 channels in Purkinje cells of S218L knockin mice was also reduced (Adams et al. 2010). In contrast, CDF of native Ca$_v$2.1 channels in both Purkinje cells (Adams et al. 2010) and Calyx of Held synaptic terminals (Inchauspe et al. 2010) of R192Q knockin mice was not significantly affected, suggesting that also the effect of FHM1 mutations on CDF likely depends on the specific Ca$_v$2.1 variant expressed in a given neuron or neuronal subcompartment.

1.5 Insights into Migraine Pathophysiology from FHM1 Knockin Mouse Models

1.5.1 Cortical Synaptic Transmission and Cortical Spreading Depression

In the cerebral cortex, excitatory synaptic transmission at pyramidal cell synapses in different cortical areas depends predominantly on P/Q-type Ca channels (Iwa-saki et al. 2000; Koester and Sakmann 2000; Rozov et al. 2001; Ali and Nelson

2006; Zaitsev et al. 2007; Tottene et al. 2009) with a notable exception at synapses between layer five pyramidal cells and burst-firing bipolar interneurons of motor cortex (Ali and Nelson 2006). Activation of the cortex following thalamic stimulation in thalamocortical slices was completely inhibited by blocking P/Q Ca channels (Llinas et al. 2007). The Ca channel pharmacology of cortical inhibitory synapses has been investigated only at fast spiking (FS) interneurons synapses; neurotransmission was found to be exclusively dependent on P/Q-type channels in many cortical areas (Zaitsev et al. 2007; Kruglikov and Rudy 2008; Tottene et al. 2009) (cf also (Sasaki et al. 2006)), but again with the exception of layer five of the motor cortex, where it was exclusively dependent on N-type (Ali and Nelson 2006).

The analysis of cortical excitatory synaptic transmission in neuronal microcultures and in brain slices from R192Q knockin mice revealed enhanced excitatory neurotransmission, due to enhanced action potential evoked Ca influx through mutant presynaptic P/Q Ca channels and enhanced probability of glutamate release at cortical pyramidal cell synapses (Tottene et al. 2009). Short-term synaptic depression during trains of action potentials was also enhanced. Neither amplitude nor frequency of miniature excitatory postsynaptic currents were altered, indicating the absence of homeostatic compensatory mechanisms at excitatory synapses onto pyramidal cells (Tottene et al. 2009). Although indirect, evidence for gain-of-function of excitatory neurotransmission has also been obtained at parallel fibers-Purkinje cell synapses in cerebellar slices of S218L knockin mice (Adams et al. 2010) and at excitatory synapses onto dorsal suprachiasmatic nucleus neurons of R192Q knockin mice (van Oosterhout et al. 2008).

In striking contrast with the enhanced glutamatergic transmission, paired recordings of fast spiking (FS) inhibitory interneurons and layer 2/3 pyramidal cells in acute cortical slices revealed that the inhibitory GABAergic transmission at FS interneuron synapses was not altered in R192Q knockin mice, despite being initiated by P/Q Ca channels (Tottene et al. 2009). The main explanation for the unaltered GABA release at FS interneuron synapses appears to be the specific expression in FS interneurons of a $Ca_v2.1$ subtype whose gating properties are little affected by the mutation (Vecchia, Tottene and Pietrobon, unpublished observations).

The investigation of experimental CSD, elicited either by electrical stimulation of the cortex in vivo or high KCl in cortical slices, revealed a lower threshold for CSD induction and an increased velocity of CSD propagation in R192Q and S218L knockin compared to WT mice (van den Maagdenberg et al. 2004, 2010; Tottene et al. 2009). Moreover, a single CSD, elicited by brief epidural application of high KCl, produced more severe and prolonged motor deficits (including hemiplegia) in FHM1 knockin mice, and, in contrast with WT mice, CSD readily propagated into the striatum (Eikermann-Haerter et al. 2009b, 2011). The much higher propensity of CSD to propagate to the striatum in FHM1 mutants compared to WT mice may explain their motor deficits and the hemiplegia typical of FHM1 aura. In agreement with the higher incidence of migraine in females, the velocity

of propagation and the frequency of CSDs, elicited by continuous epidural high KCl application, were larger in females than in males of both mutant strains; the sex difference was abrogated by ovariectomy and enhanced by orchiectomy, suggesting that female and male gonadal hormones exert reciprocal effects on CSD susceptibility (Eikermann-Haerter et al. 2009a, b).

The strength of CSD facilitation as well as the severity of the post-CSD neurological motor deficits and the propensity of CSD to propagate into subcortical structures in R192Q and S218L knockin mice were all in good correlation with the strength of the gain-of-function of the Ca$_v$2.1 channel and the severity of the clinical phenotype produced by the two FHM1 mutations (van den Maagdenberg et al. 2004, 2010; Tottene et al. 2005; Eikermann-Haerter et al. 2009b, 2011; Kors et al. 2001). Propagation of CSD to the hippocampus and thalamus and repetitive CSD events following a single CSD-inducing stimulus were observed only in S218L mutants (van den Maagdenberg et al. 2010; Eikermann-Haerter et al. 2011). These unique CSD features might account for the severe attacks with seizures, coma, and cerebral edema typical of patients with the S218L mutation. Moreover, the recent finding of increased frequency of periinfarct ischemic depolarizations during acute experimental stroke (and larger critical tissue perfusion level below which infarction ensues) in S218L knockin compared to WT mice (Eikermann-Haerter et al. 2012) suggests that the S218L mutation may lead to increased vulnerability to ischemic stroke as a consequence of increased susceptibility to ischemic depolarizations akin to CSD. The irregular firing patterns and hyperexcitability of Purkinje cells recently uncovered in S218L knockin mice likely contributes to the cerebellar ataxia produced by the S218L mutation in mice and humans (Gao et al. 2012).

Tottene et al. (2009) provided direct evidence that the gain-of-function of glutamate release at synapses onto cortical pyramidal cells may explain the facilitation of experimental CSD in FHM1 knockin mice (Tottene et al. 2009). In fact, the facilitation of CSD in acute cortical slices of R192Q knockin mice was completely eliminated (both CSD threshold and velocity became similar to those in WT slices) when glutamate release at pyramidal cell synapses was brought back to WT values by partially inhibiting P/Q channels (Tottene et al. 2009). The data are consistent with and support a model of CSD initiation in which Ca$_v$2.1-dependent release of glutamate from cortical pyramidal cell synapses and activation of NMDA receptors (and possibly postsynaptic Ca$_v$2.1 channels) play a key role in the positive feedback cycle that ignites CSD (Pietrobon 2005b; Tottene et al. 2009). This model and in general the specific requirement of Ca$_v$2.1 channels in the initiation and propagation of CSD (induced by electrical stimulation or brief pulses of high K+ in healthy tissue) are further supported by the findings that (i) after blockade of either the P/Q-type Ca channels or the NMDA receptors, CSD could not be induced in cortical slices of WT mice even with largely supra threshold depolarizing stimuli; in contrast, blockade of N- or R-type Ca channels had only a small inhibitory effect on CSD threshold and velocity of propagation (Tottene et al. 2011), and (ii) in the spontaneous mouse mutants *leaner* and *tottering*, that carry loss-of-function mutations in *cacna1a* (Pietrobon 2010), the

in vivo electrical threshold for CSD initiation was greatly increased and the CSD velocity decreased compared to WT mice (Ayata et al. 2000).

In migraineurs, CSD is not induced by experimental depolarizing stimuli, but arises "spontaneously" in response to specific triggers, that somehow create in the cortex conditions for initiation of the positive feedback cycle that overwhelms the regulatory mechanisms controlling cortical $[K^+]_o$ and ignites CSD. Insights into how this might occur have been provided by the differential effect of FHM1 mutations on cortical excitatory and inhibitory synaptic transmission (Tottene et al. 2009). This finding suggests that, very likely, the neuronal circuits that dynamically adjust the balance between excitation and inhibition during cortical activity are altered in FHM1 (Tottene et al. 2009). Functional alterations in these circuits are expected to lead to dysfunctional regulation of the cortical excitatory-inhibitory balance and hence to abnormal processing of sensory information (Monier et al. 2003; Shu et al. 2003). It has been hypothesized that this dysregulation may in certain conditions (e.g., in response to migraine triggers such as intense, prolonged sensory stimulation) lead to disruption of the excitatory-inhibitory balance and hyperactivity of cortical circuits, mainly due to excessive recurrent excitation, that may create the conditions for the initiation of "spontaneous" CSDs (e.g., by increasing the extracellular $[K^+]$ above a critical value) (Tottene et al. 2009; Vecchia and Pietrobon 2012).

Impairment of the cortical circuits that dynamically adjust the excitatory-inhibitory balance during cortical activity, due to excessive recurrent glutamatergic neurotransmission, might also underlie the abnormal regulation of interictal cortical function in some common migraine subtypes, for which there is indirect evidence consistent with enhanced cortical glutamatergic neurotransmission (Prescot et al. 2009; Siniatchkin et al. 2012) and enhanced cortico–cortical or recurrent excitatory neurotransmission (Siniatchkin et al. 2007; Wilkinson et al. 2008; Conte et al. 2010; Battista et al. 2011).

The gain-of-function effects on cortical pyramidal cell $Ca_v2.1$ channels and on cortical glutamatergic synaptic transmission derived from the functional analysis of R192Q and S218L FHM1 knockin mice provide a coherent picture of molecular and cellular mechanisms that may produce increased susceptibility to CSD and thus explain the aura symptoms in FHM1 patients (Fig. 1.2). Most likely, the conclusions are valid for all FHM1 mutations, given that the aura phenotype is common to all FHM1 patients and that eight different FHM1 mutations produce a consistent gain-of-function of single channel Ca influx through recombinant human $Ca_v2.1$ channels (see Sect 1.3). The opposite conclusions derived from the functional analysis of FHM1 mutations (including the R192Q mutation) in transfected hippocampal neurons overexpressing human $Ca_v2.1\alpha1$ subunits (i.e., loss-of-function of $Ca_v2.1$ channels with consequent decreased contribution of P/Q Ca channels to glutamatergic synaptic transmission, and unaltered synaptic strength due to compensatory mechanisms (Cao et al. 2004)) provide a picture that seems unable to explain the increased susceptibility to CSD and the aura phenotype of FHM1 patients. Thus, to be able to draw meaningful conclusions regarding