

Rodolfo Madrid · Juan Bacigalupo  
*Editors*

# TRP Channels in Sensory Transduction

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ISBN 978-3-319-18704-4

ISBN 978-3-319-18705-1 (eBook)

DOI 10.1007/978-3-319-18705-1

Library of Congress Control Number: 2015945247

Springer Cham Heidelberg New York Dordrecht London

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# Preface

In 1969, Cosens and Manning found a spontaneous mutation in *Drosophila melanogaster* that was revealed by a defect in the electroretinogram, in which the steady electrical response normally evoked by a sustained light pulse becomes transient. This impairment occurred in the receptor potential of the photoreceptor cells. For this reason, the mutant was termed Transient Receptor Potential, TRP. In 1989, Montell and Rubin cloned the gene and proposed that it encoded a novel ion channel, an idea that was confirmed in the following years by several laboratories. A second channel was also identified in the same cells by Phillips and coworkers in 1992, and was called TRP-like (TRPL), for its similarity with TRP. The first mammalian TRP channel sequence was reported independently in 1995 by the Montell and Birnbaumer laboratories, and it is known today as TRPC1, one of the seven members of the Canonical TRP (TRPC) channel subfamily, which also includes *Drosophila* TRP and TRPL. After the molecular cloning of TRPV1, the first member of the Vanilloid subfamily of TRP channels by David Julius and his group in 1997, our understanding of the molecular and cellular mechanisms underlying sensory transduction has made dramatic progress. Since then, a vast number of publications have accumulated in the literature regarding this remarkable ion channel superfamily.

The TRP superfamily is divided into seven subfamilies: TRPC (seven members in mammals with the two closely related fly channels, TRP and TRPL), TRPV (Vanilloid, six members), TRPM (Melastatin, eight members), TRPN (name derived from the fly mutant no mechanoreceptor potential C, one member), TRPA (Ankyrin, one member), TRPP (polycystic, three members) and TRPML (Mucophilin, three members) (See the phylogenetic tree of TRP channels in Chap. 1 of this book, Fig. 1.2). TRP channels are key molecular components of many physiological processes. One of the most salient features of the TRP channels is the paramount role that they play in a wide variety of sensory modalities, underlying the receptor potential in the corresponding primary sensory neurons. They participate in photo-, chemo-, thermo-, mechano- and osmoreception, pain and itching perception, and other sensory reception modalities. It is also noteworthy that, contrary to channels belonging to other superfamilies, TRP channels exhibit an impressive structural diversity, a characteristic that has contributed to complicate their classification; nev-

ertheless, the large majority of these channels possess six transmembrane domains, reminiscent of many voltage-dependent channels. Although some of these channels appear to respond mainly to a single stimulus type, many of them are polymodal, capable of being gated by stimuli of entirely unrelated nature, such as natural and artificial compounds, voltage and temperature, as is the case of the eleven thermoTRP channels that has been described. All TRPs are cation selective, many of them for monovalents, several additionally let  $\text{Ca}^{2+}$  through, while a few are strictly  $\text{Ca}^{2+}$  selective.

So far, only a small number of TRPs have been studied in some detail in intact sensory structures. An important factor for this is that they are usually confined to small and often inaccessible cellular compartments specialized in sensory transduction, such as microvilli, cilia and nerve terminals. Heterologous expression has been a powerful strategy for the study of TRP channels, mainly because it circumvents the difficulty of accessing their native location in their respective sensory cells. However, this approach must be taken with caution, because the functional properties of the channels may be largely modified when expressed in a foreign membrane, as observed in several cases (as for example *Drosophila* TRP and TRPL; Chap. 4 of this book).

This book comprises ten chapters written by experienced researchers in the field of sensory transduction, and particularly TRP channels. The origin of the present book was a symposium on TRP channels and sensory transduction, organized by the Editors for the First Meeting of FALAN (Federation of Neuroscience Societies in Latin America, the Caribbean and the Iberian Peninsula) in Cancún, México, in 2012. This symposium interested Springer NY, which kindly offered us to carry out this book.

The aim of this book is to offer up-to-date discussions regarding some iconic members of this highly interesting ion channel superfamily, which additionally are among those that are best characterized. Rather than extensive and exhaustive, the chapters attempt to be comprehensive to a non-specialized audience, as well as informative to those in the field. Inevitably, for instructive purposes, a small number of technical issues treated in a formal manner were included in some chapters. We felt that these formalisms were necessary for the comprehension of the thermodynamics and dynamic modeling of thermoTRPs.

We would like to thank the authors for their excellent contributions that made this book possible. The editors also thank Simina Calin and Portia Wong, Editors of Springer NY, for efficient guidance, support and excellent editorial advice.

We hope that the book will be found useful for Neurobiology students, teachers and specialist in sensory transduction. Enjoy to this TRiP to the Senses.

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# Contents

<b>1 Biophysical and Molecular Features of Thermosensitive TRP Channels Involved in Sensory Transduction</b> .....	1
Gonzalo Ferreira, Natalia Raddatz, Yenisleidy Lorenzo, Carlos González and Ramón Latorre	
<b>2 Pharmacology of TRP Channels</b> .....	41
Asia Fernández-Carvajal, Gregorio Fernández-Ballester, Rosario González-Muñiz and Antonio Ferrer-Montiel	
<b>3 Modulation of TRP Channels by N-glycosylation and Phosphorylation</b> .....	73
María Pertusa and Rodolfo Madrid	
<b>4 TRP Channels in Visual Transduction</b> .....	97
Juan Bacigalupo, Ricardo Delgado, Yorika Muñoz and Peter O’Day	
<b>5 TRP Channels in Transduction for Responses to Odorants and Pheromones</b> .....	111
Diego Restrepo, Rona Delay, Weihong Lin, Fabián López and Juan Bacigalupo	
<b>6 TRP Channels as Targets for Modulation of Taste Transduction</b> .....	127
Karel Talavera	
<b>7 TRP Channels and Mechanical Transduction</b> .....	141
Ana Gomis	
<b>8 TRP Channels in the Sensation of Heat</b> .....	165
Chun-Hsiang Tan and Peter A. McNaughton	

**9 TRP Channels in Cold Transduction** ..... 185  
Alejandro González, Gonzalo Ugarte, Ricardo Piña, María  
Pertusa and Rodolfo Madrid

**10 Mathematical Modeling of TRPM8 and the Cold  
Thermoreceptors** ..... 209  
Erick Olivares and Patricio Orio

**Index** ..... 225



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# Chapter 1

## Biophysical and Molecular Features of Thermosensitive TRP Channels Involved in Sensory Transduction

Gonzalo Ferreira, Natalia Raddatz, Yenisleidy Lorenzo,  
Carlos González and Ramón Latorre

**Abstract** Temperature is one of the physical variables that cells and biological organisms constantly monitor to achieve homeostasis and maintain chemical reactions at a suitable speed for the living environment to which they are adapted. In order to monitor and maintain temperature on a constant basis, thermosensitive molecules were selected during evolution. One of the most remarkable sets of molecules acting as sensors is constituted by thermosensitive transient receptor potential channels (thermoTRP channels). TRP channels are a superfamily of non-selective tetrameric cation channels closely related to the classic superfamily of voltage-gated channels, having a set of distinctive sequence elements in common, while acting as polymodal receptors. This latter ability is what makes them suitable for integrating many kinds of signals in different cells, ranging from chemical to physical stimulation (*i.e.*: temperature-, mechano- and chemo-sensitivity). These channels act as allosteric proteins modifying sensitivity to one stimulus in the presence of another, and thus allowing the integration of many different signaling processes that are critical for sensing the extracellular and intracellular environment and for maintaining homeostasis. This ability has made them vital for life support. Several subfamilies of TRP channels have been described. From these subfamilies, some types of channels have been distinguished as being temperature-sensitive, such as TRPV1–4, TRPM 2–5/8, TRPA1 and TRPC5. In this chapter, thermosensitivity will be defined. Then, we will describe the thermosensitive molecules identified so far, focusing our analysis on ion channels, particularly on thermosensitive TRP channels involved in sensory transduction. Their gating and permeation properties and

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© Springer International Publishing Switzerland 2015  
R. Madrid, J. Bacigalupo (eds.), *TRP Channels in Sensory Transduction*,  
DOI 10.1007/978-3-319-18705-1\_1

gating modifiers shall be at the center of the discussion so as to place them in the context of ion channels and life evolution.

**Keywords** ThermoTRP channels · Biophysics of TRP channels · Sensory transduction

## 1.1 Thermosensitivity in Organisms and Across Species

Thermosensitivity is the property of inorganic and organic materials to respond to changes in heat or temperature faster than the average speed of response observed in comparable kinds of substances found in nature. In this chapter, we will focus on the thermosensitivity found in living organic materials, particularly in organisms, cells and molecules.

Thermal variation is among the oldest types of stress that biological beings have been subjected to, and adaptation responses to it usually involve certain pathways conserved through evolution. Regarding organisms, thermosensitivity can be found in eukaryotes and prokaryotes. In complex eukaryotes, total and local body temperature control has been measured in birds and mammals, where it has been found that total body thermosensitivity generally varies between  $-4$  and  $-12$  W/(kg°C) (Mercer and Simon 1984). In reptiles and fish, it has been reported that sexual determination is especially thermosensitive (Baroiller and D’Cotta 2001; Pieau et al. 2001). Whole body temperature adjustment, in particular, has been related to neurons of the central nervous system and sensing in peripheral receptors (Hori and Katafuchi 1998). Phenomena related to cold and heat shock genes that are expressed with sudden changes in temperature have been extensively reported and studied in prokaryotes (Hurme and Rhen 1998). It is interesting that, in general, such responses seen in prokaryotes are conserved among more complex organisms such as eukaryotes (Al-Fageeh and Smales 2006).

### 1.1.1 A Biophysical Description of Thermosensitivity

Thermosensitivity for whole organisms or materials is physically expressed by specific heat units. Usually, the thermosensitivity of molecules or biological processes is measured by its  $Q_{10}$  (or temperature coefficient), which is a measure of the rate of change in biological or chemical systems resulting from increasing the temperature by  $10^\circ\text{C}$ . In general, it is calculated as follows:

$$Q_{10} = \left( \frac{k_1}{k_2} \right)^{\frac{10}{T_1 - T_2}} \quad (1.1)$$

where  $k_1$  and  $k_2$  are the rate constants of the reaction proportional to the rates of the process at two different temperatures ( $T_1$  and  $T_2$ ), normally using the Kelvin

scale.  $Q_{10}$  for most biological systems and molecules measured in physiological conditions is generally between 2 and 3, whereas for thermosensitive molecules or biological processes it is usually more than 5 (Nelson et al. 2008). However,  $Q_{10}$  has several limitations in measuring thermosensitivity since it is quite dependent on the ratio of temperatures used to measure it.

A more accurate description of thermosensitivity and the dependence of reaction rates on temperature are related to Arrhenius energy of activation, which gives us a relationship between the kinetic constants of chemical reactions and the energy activation:

$$k_x = Ae^{\frac{-Ea}{RT}} \quad (1.2)$$

where  $k_x$  is the velocity constant,  $A$  is the prefactor, being a constant related to the frequency of molecular collisions,  $Ea$  is the energy of activation,  $R$  is the gas constant and  $T$  is absolute temperature.

From Eqs. (1.1) and (1.2), we have:

$$Ea = -RT_1 \left( \frac{T_1 + 10}{10} \right) \ln Q_{10} \quad (1.3)$$

Thus, a thermosensitive molecule with a  $Q_{10} \sim 10$  implies an  $Ea \sim 40$  cal/mol at a temperature of 298 K.

In transition state theory, activation energy by Arrhenius can be somehow related to the following equation:

$$k = -\kappa \frac{k_B T}{h} e^{-\frac{\Delta G^*}{RT}} \quad (1.4)$$

where  $\kappa$  is the transmission factor,  $\frac{k_B T}{h}$  is the frequency factor (also called the pre-exponential factor) which has the units of  $s^{-1}$  and  $\Delta G^*$  is the standard Gibbs energy of activation. As it can be seen, it is similar to Eq. 1.2 derived from Arrhenius statements for the whole reaction, with  $A = \kappa \frac{k_B T}{h}$ . Taking into account that:

$$\Delta G^* = \Delta H^* - T\Delta S^* \quad (1.5)$$

where  $\Delta H^*$  is the enthalpy change,  $T$  is the absolute temperature and  $\Delta S^*$  is the entropy change, we reach the following relationship between the energy of activation  $Ea$  from Arrhenius, the transition state theory (Eq. 1.5):

$$Ea = \Delta H^* - T\Delta S^* \quad (1.6)$$

Therefore, a molecule will be more thermosensitive when the enthalpy differences to reach the active state at two given temperatures are higher. Keeping in mind that enthalpy and entropy are functions of temperature as well, a general expression

for Gibbs free energy at a given condition would be as follows (Digel et al. 2008; Nelson et al. 2008):

$$\Delta G = \Delta H_{(TR)} - T\Delta S_{(TR)} + \Delta C_p(T - TR) + \Delta\Delta C_p \ln\left(\frac{T}{TR}\right) \quad (1.7)$$

where  $TR$  is an arbitrary reference temperature and  $\Delta C_p$  is the heat capacity change for a particular protein or molecule between temperature  $T$  and the reference temperature  $TR$ . This equation allows us to know the limits of protein stability when there are temperature changes (heat or cold).

The temperature dependence of  $\Delta H$  and  $\Delta S$  is usually parabolic. This will be quite evident when  $C_p$  is high, meaning that in those cases we might find denaturation by hot and cold temperatures as extreme examples of variations in molecule conformation due to temperature. Most interactions that yield particular conformations at physiological temperatures and that stabilize proteins in their native environments are weak non-covalent bonds (Petsko and Ringe 2004).

Thermosensitive molecules could also react to other physical parameters (i.e., voltage, mechanics, degrees of hydration, etc.). Allosteric interactions between the variables that control protein conformations would arise quite easily. Hence, any attempt to construct gating kinetic models for thermosensitive channels should be allosteric (Matta and Ahern 2007; Latorre et al. 2007a, b; Baez-Nieto et al. 2011; Jara-Oseguera and Islas 2013).

### 1.1.2 Thermosensitive Molecules and Sensory Transduction

Temperature can promote dramatic changes in cells ranging from membrane fluidity, DNA-RNA processes and structure-catalysis from enzymes (Digel et al. 2008). All these elements and their modifications due to heat/temperature can serve as molecular sensors for thermosensitivity. Regarding *membranes*, phase-transition changes occur with sudden transitions in temperature, which might influence gene expression (Vigh et al. 1998). Changes in membrane fluidity due to temperature can trigger the expression of enzymes that would modify molecular composition.

Alterations in *nucleic acids* seem to constitute a more efficient biomolecular sensor for temperature changes and gene expression. The 3D architecture of RNA has been recently described as a critical sensor event, being part of what is known as *riboswitches* (Mandal and Breaker 2004). There are two regions in this complex ribosomal RNA constituent, namely the aptamer and expression platforms. The aptamer platform has been described to sense changes in molecules promoted by heat/temperature and, hence, to regulate gene expression at the translation level. Regarding DNA, the supercoiling level and topoisomerases appear to be critically sensitive to temperature (Tse-Dinh et al. 1997).

Ultimately, the critical biomolecular targets for temperature control and sensing are *proteins*. In *E. coli*, chemoreceptor protein systems such as *Tar*, *Tsr*, *Trg* and *Tap*

have been described. What is interesting is that they share a significant degree of homology in spite of the different ranges in thermal sensitivity they all exhibit (Nara et al. 1991). In complex organisms, specialized ion channels are able to transduce heat/cold into channel opening. Sensory cells and several tissues express thermosensitive molecules, from which thermoTRP channels are a clear example and are directly or indirectly involved in the homeostasis of temperature as a physiological phenomenon (Digel et al. 2008).

Here, we will discuss these concepts in molecules, with a particular emphasis on ion channels as specialized sensors and effectors that have evolved to sense and control heat and temperature in living organisms. In particular, we will center our discussion on thermosensitive TRP channels, which play a critical role in sensory thermotransduction-

## 1.2 Thermosensitive Channels and Thermal Transduction

Our skin is a physical barrier that, among other functions, enables us to discriminate temperatures ranging from extreme cold (about  $-10^{\circ}\text{C}$ ) to extreme heat (about  $60^{\circ}\text{C}$ ). The neurons that allow us to sense this wide range of temperatures are located in the trigeminal ganglia (Walder et al. 2002) innervating the face and head and in the dorsal root ganglia (DRG) for the rest of the body. The process of thermosensation begins with specialized proteins which are expressed in the free nerve endings of the afferent fibers innervating the skin and respond to external temperature changes.

All proteins including ion channels and indeed all biological or chemical processes are sensitive to temperature with a  $Q_{10}$  value of  $\sim 2$  (Hille 2001). However, thermo-TRP receptor channels present an exceptionally high  $Q_{10}$  value  $\geq 20$ . To date, at least 10 TRP channels have been observed to exhibit steep temperatures, namely TRPV1, TRPV2, TRPV3, TRPV4, TRPM2, TRPM3, TRPM4 and TRPM5, which are activated upon warming, and TRPM8, TRPA1 and TRPC5, which are activated upon cooling (Vriens et al.; Caterina et al. 1997, 1999; McKemy 2002; Peier 2002a, b; Smith et al. 2002; Watanabe et al. 2002b; Xu et al. 2002; Chung et al. 2003; Story et al. 2003; Talavera et al. 2005; Togashi et al. 2006; Karashima et al. 2009; Zimmermann et al. 2011). They show different activation thresholds covering the range of thermal oscillations to which living organisms are usually exposed. For example, TRPV1 is activated by temperatures in the noxious range ( $>42^{\circ}\text{C}$ ), whereas TRPM8 in neurons is mainly closed at temperatures  $>33^{\circ}\text{C}$  and is primarily open at lower temperatures, allowing sodium and calcium ions to flow into the nerve terminals leading to depolarization, which in turn triggers action potential firing (Babes 2009).

Madrid and coworkers (Madrid et al. 2009), found that different expression ratios between TRPM8 and Kv1 tune cold sensitivity in cold thermoreceptor neurons. They pointed out that low cold thresholds can be produced by high TRPM8 expression levels and low or absent expression of  $I_{\text{KD}}$  a (a current dependent on Shaker-like Kv1.1–1.2 channels that act as an excitability brake). In contrast, specific high-

threshold cold neurons would result in low TRPM8 levels and high levels of  $I_{KD}$  expression (Madrid et al. 2009). Thus, apparently TRPM8 and  $I_{KD}$  play a critical role in determining the thermal excitability of cold-sensitive neurons (See Chap. 9 by Gonzalez et al. in this book).

In addition to TRPs, other ion channels participate in thermal transduction. Leak or background potassium channels, namely TREK-1, TREK-2 and TRAAK, members of the two-pore domain  $K^+$  (K2P) channel family, have been shown to be sensitive to temperature in different expression systems such as *Xenopus oocytes* and COS cells (Kang 2005). TREK-1 has shown low activity at room temperature, which increases with rising temperatures and reaches maximum activation at 37°C (Maingret et al. 2000). This suggests that TREK-1 is active at body temperature, thus contributing significantly to background  $K^+$  conductance in physiological conditions. Moreover, TREK-2 and TRAAK whole-cell currents show a ~20-fold increase when bath temperature rises from 24 to 42°C, with activation thresholds of 25 and 31°C, respectively (Kang et al. 2005). These channels tend to hyperpolarize neurons, closing rapidly and reversibly with small temperature reductions.

Finally, hyperpolarization-activated nonselective cationic channels of the HCN family, specifically HCN1, play an important role in determining the firing patterns of cold receptors in mammals, hence contributing to the ability of neurons to respond to stimuli with high frequency bursts (Babes 2009). Although it does not seem to be involved in responses to acute cooling, mice lacking the *hcn1* gene show altered cold perceptions (Orio et al. 2009). Accordingly, an ample group of thermo-sensitive ion channels has been discovered but further studies are still required to better understand the global phenomenon of thermo-transduction.

## 1.3 TRP Channels

### 1.3.1 A Brief History of TRP Channels

Transient Receptor Potential genes (*trp*) were first described in the late 1960s in the fruit fly *Drosophila melanogaster* (Cosens and Manning 1969). The name is derived from a spontaneously occurring mutation in the *Drosophila* fly, resulting in visual transduction defects. The eyes of these mutant flies elicit a transient electric response to continuous light instead of a continuous response, as it is the case of the wild type flies. Twenty years later, it was reported that the *trp* gene encodes a large protein, namely the TRP channel, consisting of 1275 amino acid residues displaying eight putative transmembrane (TM) segments (Montell and Rubin 1989). In addition, the amino acid sequence did not present clear homologies with any other known ion channel.

Just 3 years later, Phillips et al. (1992) found a new gene that was named TRP-like (*trpl*). The protein encoded by *trpl* showed structural homologies with *Drosophila* TRP and also with the superfamily of voltage-gated channel genes (Phillips et al. 1992). In the same year, Hardie and Minke demonstrated that the TRP gene

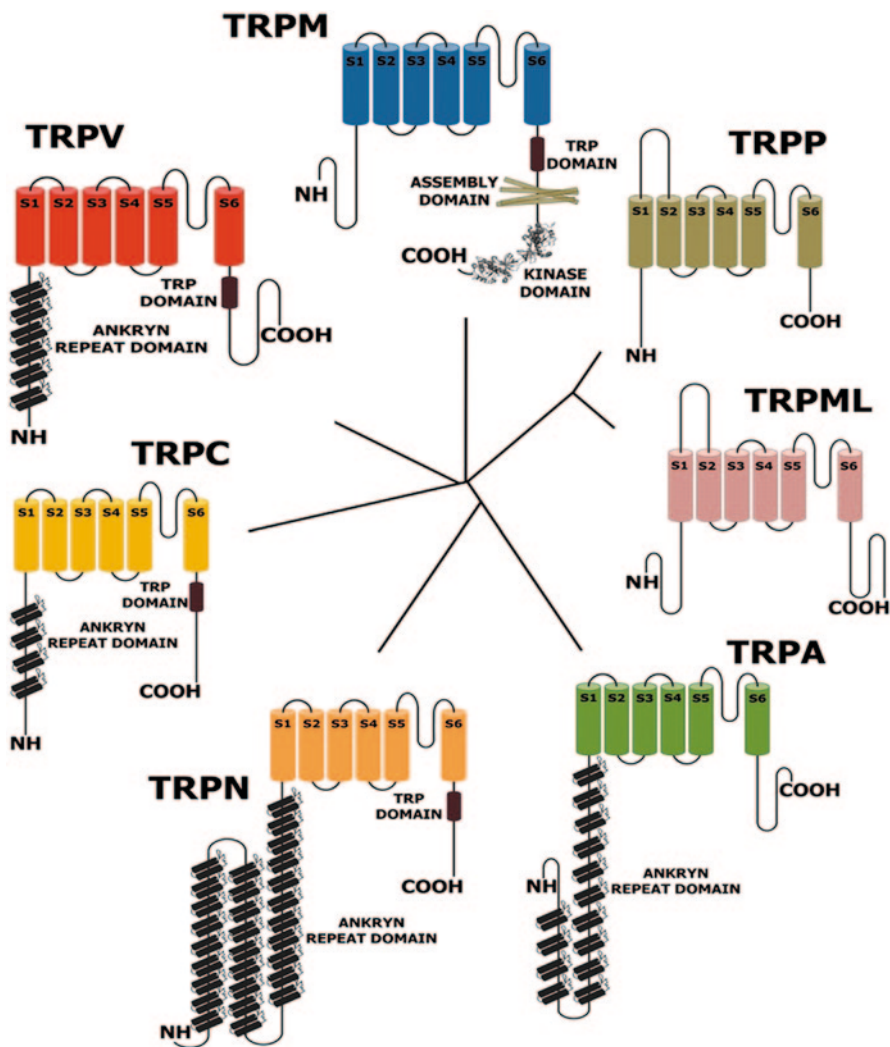


coded for a highly  $\text{Ca}^{2+}$  permeable channel (Hardie and Minke 1992). In 1995, Petersen revealed that TRP related proteins were also present in vertebrates, such as *Xenopus laevis* and mice cDNA libraries (Petersen et al. 1995), while Wes and Zhu reported the full sequence of the first human homologue, TRPC1, which shares 37% identity with the *Drosophila trp* gene (Wes et al. 1995; Zhu et al. 1995). Today, we can list more than 100 channels within this family in both invertebrates and vertebrates (Nilius and Owsianik 2011).

### 1.3.2 *The Superfamily of TRP Channels and Thermosensitive TRP Channels*

The TRP superfamily comprises a large number of ion channels with diverse functions, serving as cellular sensors of physical and chemical stimuli such as light, temperature, voltage, touch, pain, osmolarity, pheromones, phosphatidylinositol 4,5-bisphosphate ( $\text{PIP}_2$ ),  $\text{Ca}^{2+}$ , cyclic nucleotides, among others (Clapham et al. 2001; Clapham 2003; Ramsey et al. 2006). For example, TRPV1 responds to heat, capsaicin and low pH, while TRPM8 is activated by menthol and cold. Hence, TRP channels are involved in the classic sensory transduction processes of multicellular organisms, such as vision, hearing, smell, taste, touch, temperature, and pain sensation. The division into subfamilies on the basis of amino acid sequences and structural similarities does not provide a functional classification for TRP proteins. For example, some members of the TRPV subfamily are involved in thermal and nociceptive sensing (TRPV1–4), whereas others are insensible to temperature but are highly selective to  $\text{Ca}^{2+}$ , thus participating in the epithelial  $\text{Ca}^{2+}$  transport (TRPV5–6).

Structurally, all members of the TRP family include six putative transmembrane domains, cytosolic N- and C-terminal tails and an ion pore between transmembrane domains 5 and 6 (Montell 2005) (Fig. 1.1). According to their amino acid sequence homology, TRP channels are grouped into seven subfamilies which, in turn, are classified into groups 1 (TRPC, TRPV, TRPM, TRPA, and TRPN) and 2 (TRPPP and TRPML) based on sequence and topological differences (Montell et al. 2002a). Group 1 of TRPs shows the strongest sequence homology with the founding member of the superfamily, *Drosophila* TRP (Montell and Rubin 1989). Group 2, on the other hand, includes distant relatives that share sequence homologies over the transmembrane segments and contain a large loop separating the first two transmembrane domains. An eighth subfamily (TRPY, where Y stands for *yeast* TRPs) is distantly related to Group 1 and Group 2 (Palmer et al. 2001; Denis and Cyert 2002).



**Fig. 1.1** TRP channel subfamilies and schematic structure of their major subunits. TRP channels present four subunits containing six transmembrane (*TM*) segments (S1–S6). Ankyrin domains (number variable) are present in the N-terminal region of TRP channels in Group 1 subfamilies. The TRP domain (well-conserved region) and protein kinase domain are present in the C terminus region

### 1.3.2.1 Group Subfamilies

#### Group 1 Subfamilies

**TRPC (classic)** TRPC channels are ubiquitously expressed among cell types and mediate signals in response to phospholipase C (PLC)-coupled receptors (Clapham 2003; Montell 2005). They have four N-terminal ankyrin-like repeat domains

(ARD), a TRP box, a homer-binding region, a calmodulin  $IP_3$ -receptor binding site (CIRB) and a PDZ domain in TRPC4-5. The first mammalian homologues of *Drosophila* TRPs were referred to as TRPC1, TRPC2, and TRPC3, which share 40% homology (Wes et al. 1995; Zhu et al. 1995). Since then, seven mammalian TRPC proteins (TRPC1-7) have been described (Montell et al. 2002b). Their subunits can assemble into homomeric and heteromeric channels (Goel et al. 2002; Hofmann et al. 2002; Strubing et al. 2003; Schaefer 2005), and they have been proposed to be activated by a variety of signals including  $Ca^{2+}$  store depletion, membrane lipids and vesicular insertion into the plasma membrane.

**TRPV (vanilloid)** The TRPV subfamily of ion channels contains three ankyrin repeats, a TRP box and calmodulin binding site. It consists of six members, of which TRPV1-4 are temperature-sensitive non-selective cation channels, and TRPV5-6 are channels insensitive to temperature and the most  $Ca^{2+}$  selective ( $P_{Ca^{2+}}/P_{Na^{+}} > 100$ ) channels among mammalian TRPs (Montell 2005). Mammalian channels TRPV1, TRPV2 and TRPV4 are also involved in sensing osmolarity changes and mechanical stimuli (Kim et al. 2003; Gong et al. 2004).

**TRPM (melastatin-related)** TRPM channels share four regions of high homology (TRPM homology regions—MHRs) in their N-terminal cytoplasmic segment. The TRPM subfamily contains eight members (TRPM1-8) that share a cytoplasmic coiled-coil domain and a well-conserved TRP box, but have diverse C-terminal domains with structures that are important in controlling the ion channel activation mechanism. TRPM6/7 and TRPM2 are designated as chanzymes, since their C-terminal contains catalytic domains that show protein kinase activities and ADP-ribose pyrophosphatase activities, respectively (Nadler et al. 2001; Perraud et al. 2001; Runnels et al. 2001; Sano et al. 2001; Schlingmann et al. 2002; Walder et al. 2002). TRPM2-5 are heat activated, while TRPM8 is activated by cold temperature. In contrast to all other TRPs, both TRPM4 and 5 form ion channels permeable to monovalent cations but not to  $Ca^{2+}$  (Launay et al. 2002; Hofmann et al. 2003; Nilius et al. 2003). TRPM3 has been reported to be activated by noxious heat (Vriens et al. 2011) and by reductions in extracellular osmolarity in HEK-293 cells (Grimm et al. 2003).

**TRPA (ankyrin)** This family includes only one member present in humans and mammals, namely TRPA1. The N-terminal region of the TRPA1 protein comprises 17 ankyrin repeats, which are common structural motifs that mediate protein-protein interactions (Gaudet 2008). It is expressed in primary somatosensory neurons (30% co-expressed with TRPV1), keratinocytes, astrocytes, vascular smooth muscle and endothelium of the cardiovascular system, gastrointestinal tract, respiratory system, pancreas, inner ear and odontoblasts. It is activated by different chemical compounds, including the psychoactive components from marijuana (i.e., delta-9-tetrahydrocannabinol), environmental irritants, products from inflammatory responses and pungent compounds of wasabi, garlic (allicin), cinnamon oil (cinnamaldehyde), mustard oils (allyl isothiocyanate) and tear gas (acrolein) (Jordt et al. 2004). TRPA1 has been reported to be a cold receptor involved in the detection of noxious cold and hypersensitivity to cold (Story et al. 2003; del Camino et al. 2010) (see below).

**TRPN (no mechanoreceptor potential C [NOMP-C homologues])** This family has a single member that can be found in worms, flies, and zebrafish (Littleton and Ganetzky 2000; Walker et al. 2000; Sidi et al. 2003). Mammals do not encode any TRPN homologues. TRPNs lack the TRP domain, but they do have the conserved TRP-box. Apparently they have a role in mechanotransduction (Walker et al. 2000).

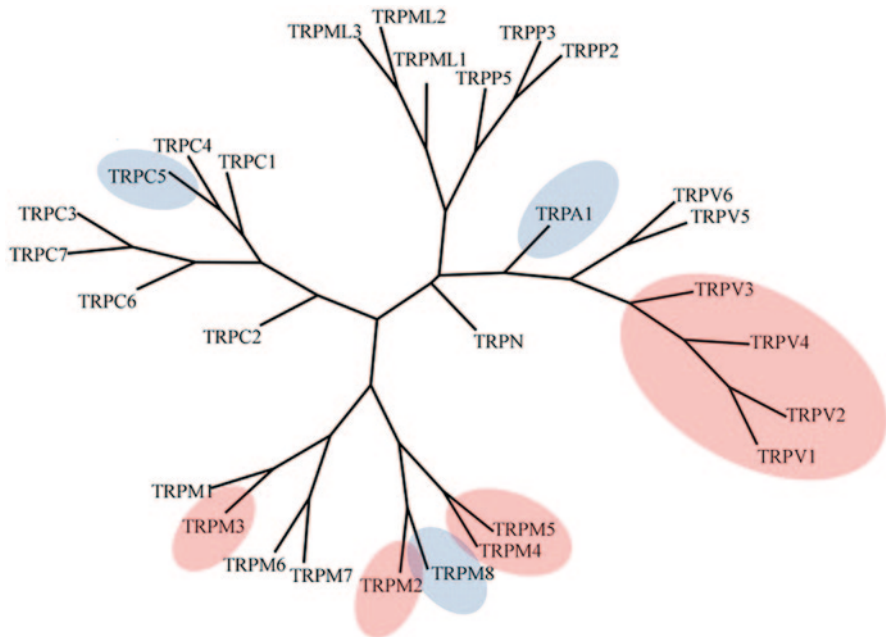
### Group 2 Subfamilies

**TRPP (polycystin)** The TRPP subfamily is formed by polycystic kidney disease (PKD) proteins or polycystins. It is formed by three members, namely TRPP2 (also known as PKD2), TRPP3 (or PKD2L1), and TRPP5 (or PKD2L2). TRPP channels carry an EF-hand motif (structural domain in proteins that bind  $\text{Ca}^{2+}$ ) and an ER retention signal. TRPP channels are present in both motile and primary cilia, and may function to sense fluid flow, osmolarity, and mechanical stretch (Stayner and Zhou 2001; Nauli et al. 2003; Venkatachalam and Montell 2007).

**TRPML (mucolipin)** The TRPML subfamily is defined by a human protein (i.e., TRPML1) that forms a non-selective  $\text{Ca}^{2+}$  channel (Bargal et al. 2000; Bassi et al. 2000; Sun et al. 2000). Mutations in this protein are responsible for the lysosomal storage disorder *mucopolipidosis IV*, which is characterized by severe neurodegeneration. In addition to TRPML1, mammals encode two other closely related proteins referred to as TRPML2 and TRPML3. TRPML1 and TRPML2 are expressed in the lysosomal membrane, while TRPML3 resides predominantly in the endoplasmic reticulum (ER) membrane in cultured cells (Manzoni et al. 2004; Kiselyov et al. 2005; Venkatachalam et al. 2006).

### 1.3.2.2 ThermoTRP Channels

Cloning and characterizations of a temperature-sensitive subclass of the TRP channels (i.e., thermoTRPs) have shed light on our understanding of the molecular basis of thermal sensation. As we mentioned before, to date, ten TRP channels have been reported to be directly activated at specific temperature ranges covering from noxious heat to painful cold. TRPV1, TRPV2, TRPV3 and TRPV4 are heat-activated, whereas TRPM8, TRPA1 and TRPC5 are activated by cold (McKemy et al. 2002; Peier et al. 2002a; Story et al. 2003; Talavera et al. 2005; Dhaka et al. 2006; Huang et al. 2006; Togashi et al. 2006; Zimmermann et al. 2011) (see Fig. 1.2). TRPM2, TRPM3, TRPM4 and TRPM5 also show temperature sensitivity but are not usually included in the thermo-TRP family because they are not expressed in primary somatosensory neurons and need adequate concentrations of intracellular  $\text{Ca}^{2+}$  to exhibit temperature sensitivity. Nevertheless, the fact that they can be temperature sensitive with an elevated  $Q_{10}$  in particular conditions has led us to include them in this chapter.



**Fig. 1.2** Phylogenetic tree of TRP channels. ThermoTRP channels are indicated according to thermosensitivity: cold (*blue*) and heat (Zaritsky et al. 2001). Dendrogram was based on full length sequence comparisons of human TRP channel protein using the UniProt services website: TRPV1 (Q8NER1); TRPV2 (Q9Y5S1); TRPV3 (Q8NET8); TRPV4 (Q9HBA0); TRPV5 (Q9NQA5); TRPV6 (Q9H1D0); TRPM1 (Q7Z4N2); TRPM2 (O94759); TRPM3 (A2A3F7); TRPM4 (Q8TD43); TRPM5 (Q9NZQ8); TRPM6 (Q9BX84); TRPM7 (Q96QT4); TRPM8 (Q7Z2W7); TRPC1 (P48995); TRPC3 (Q13507); TRPC4 (Q9UBN4); TRPC5 (Q9UL62); TRPC6 (Q9Y210); TRPC7 (Q9HCX4); TRPML1 (Q9GZU1); TRPML2 (Q8IZK6); TRPML3 (Q8TDD5); TRPP2 (Q13563); TRPP3 (Q9P0L9); TRPP5 (Q9NZM6); TRPA1 (O75762). TRPC2 is a pseudogene in humans and TRPNs are not present in mammals; therefore we used mouse TRPC2 (Q9R244) and zebrafish TRPN (Q7T1G6)

**TRPV1** This TRP channel was first characterized by McNaughton and his group (Cesare and McNaughton 1996), and was subsequently cloned by the group led by David Julius (Caterina et al. 1997). It is a non-selective cation permeable channel activated by a wide range of stimuli including capsaicin and low pH (<6), and by physical factors such as heat and membrane depolarization (Caterina et al. 1997; Tominaga et al. 1998; Voets et al. 2004). Activation threshold by heat is in the range of 41–43 °C, likely reaching a maximum over 50 °C. This channel can be directly activated by ethanol (Trevisani et al. 2002), several endogenous lipids (i.e., endocannabinoid, anandamide and N-arachidonoyl dopamine) (Ross 2003; De Petrocellis et al. 2004), metabolic products of lipoxygenase (Hwang et al. 2000), topical analgesics such as camphor (Xu et al. 2005), and pungent compounds present in black pepper (piperine) (McNamara et al. 2005) and garlic (allicin) (Macpherson et al. 2005; Salazar et al. 2008). Consequently, TRPV1 is viewed as a signaling integrator for many noxious stimuli.

**TRPV2** Although this channel shares 50% of sequence homology with TRPV1, it is insensitive to capsaicin, low pH and responds to higher temperatures with an activation threshold of 52 °C (Caterina et al. 1999). TRPV2 is activated in vitro by three different types of physical stimuli: heat, osmotic stress and mechanical stretch (Caterina et al. 1999; Muraki et al. 2003). It is also activated by chemicals such as 2-aminoethoxydiphenyl borate (2-APB) (Hu et al. 2004) and by cannabidiol, a natural component of the marijuana plant (Qin et al. 2008). The physiological role of TRPV2 in thermal nociception remains unknown because TRPV2<sup>-/-</sup> mice showed no evident deficits in thermal sensation (Park et al. 2011).

**TRPV3** This ion channel is implicated in the perception of warmth in the skin with an activation threshold of 34–38 °C (Peier et al. 2002b; Smith et al. 2002; Xu et al. 2002). TRPV3 is activated by 2-APB (Chung et al. 2004) and by various natural compounds such as carvacrol, thymol and eugenol, which are present in essential oils from herbs and spices such as *Thymus vulgaris*, *Origanum syriacum*, and *Caryophylliflos* (Moqrich et al. 2005; Xu et al. 2006). TRPV3<sup>-/-</sup> mice showed some deficits in sensing hot temperatures in the innocuous to noxious range (Moqrich et al. 2005).

**TRPV4** TRPV4 is a non-selective cation channel that shares ~ 40% amino acid identity with TRPV1 (Liedtke et al. 2000; Strotmann et al. 2000; Delany et al. 2001). This thermo-TRP is activated by moderate temperatures (27–34 °C) (Watanabe et al. 2002b), by hypotonic solutions, suggesting that it serves as a sensor for osmolarity or mechanical stretch (Liedtke et al. 2000; Strotmann et al. 2000; Wissenbach et al. 2000; Delany et al. 2001; Alessandri-Haber et al. 2005), and by phorbol esters (Watanabe et al. 2002a). Knockout studies suggest that TRPV4 also plays a role in thermal hyperalgesia (Todaka et al. 2004).

**TRPM2** TRPM2 channels are Ca<sup>2+</sup>-selective TRP channels, modulated by ADP-ribose, intracellular Ca<sup>2+</sup> and oxidative stress. In the presence of ADP-ribose and intracellular Ca<sup>2+</sup>, they are heat-activated by warm temperatures. They are expressed mainly in the brain, pancreas and immune cells. As for most of the TRP channels, they also have several modulating pathways in addition to those just mentioned (Eisfeld and Luckhoff 2007).

**TRPM3** TRPM3 channels are non-selective cation channels permeable to Ca<sup>2+</sup>, which were classically described as osmo- and volume-regulated TRP channels, denoting properties in common with TRPV4 channels. They are mostly expressed in the kidney, neurons and eyes, and have several splice variants in the N and C-terminus plus the S1-S6 region (some of them including the pore region) (Grimm et al. 2003; Oberwinkler and Philipp 2007). These channels are activated by noxious heat (> 40 °C) and are also strongly associated and modulated by steroids (pregnenolone sulfate), acting as membrane receptors for these lipids (Wagner et al. 2008).

**TRPM4 and TRPM5** Both are non-selective Ca<sup>2+</sup>-activated cation channels. They also share 50% of their sequences, being modulated by similar mechanisms. Sev-



eral splice variants have been identified for TRPM4 channels (a, b and c), being TRPM4b the most widespread. They are both heat-activated in a warm temperature range and in the presence of intracellular  $\text{Ca}^{2+}$ . Just like most TRP channels, their function is modulated by a wide range of extra and intracellular ligands. Whereas TRPM4s are widely expressed throughout the organism, TRPM5s appear to be confined mostly to taste receptors (Talavera et al. 2005; Guinamard et al. 2011).

**TRPM8** TRPM8 was the first TRP channels found to sense cold temperatures (McKemy 2002; Peier 2002a). Activation is measurable at  $\sim 33^\circ\text{C}$  in sensory neurons, with currents increasing in magnitude at lower temperatures (McKemy et al. 2002; Peier 2002a; Brauchi 2004; Brauchi et al. 2004; Madrid et al. 2006; Bautista et al. 2007; Malkia et al. 2007). Furthermore, this channel is activated by voltage (Brauchi 2004; Voets et al. 2004) and chemical agonists that mimic cooling, such as menthol, eucalyptol or icilin (McKemy 2002; Peier 2002a; Chuang 2004; Boddington et al. 2007). TRPM8 is also activated by the regulatory lipid  $\text{PIP}_2$  (Liu and Qin 2005; Rohacs et al. 2005). In vivo studies of TRPM8<sup>-/-</sup> mice showed deficits in reacting to cool temperatures, indicating that this channel is a major sensor of peripheral innocuous coolness (Bautista et al. 2007; Colburn et al. 2007; Dhaka et al. 2007).

**TRPA1** TRPA1 is activated within the noxious cold range of temperatures ( $\sim \leq 17^\circ\text{C}$ ), suggesting that it may be responsible for detecting painful levels of cold (Story et al. 2003). The active constituents of mustard oil (allyl isothionate) and garlic (allicin) robustly activate TRPA1 currents (Bandell et al. 2004; Jordt et al. 2004; Macpherson et al. 2005). TRPA1 is directly activated by  $\text{Ca}^{2+}$  binding to an EF-hand domain located within its intracellular N-terminal region. This suggested that apparent cold sensitivity may not result from direct temperature-sensitive gating, but instead from a cold induced increase in intracellular  $\text{Ca}^{2+}$  (Doerner et al. 2007; Zurborg et al. 2007). Furthermore, analyses of TRPA1<sup>-/-</sup> mice has yielded conflicting data, with one group reporting no deficits in acute cold sensing, while a second group observing reduced cold sensitivity in female but not in male mice, and a third group showing cold deficits after prolonged exposure to cold (Bautista et al. 2006; Kwan et al. 2006; Karashima et al. 2009). However, these apparent conflicting results are a consequence of species differences. Thus, cold activates rat and mouse TRPA1 but not human or rhesus monkey TRPA1 (Chen et al. 2013).

**TRPC5** TRPC5 is cold-sensitive in the temperature range of  $37\text{--}25^\circ\text{C}$  ( $Q_{10} \sim 10$ ). It is present in mouse and human sensory neurons of dorsal root ganglia and in the dorsal lamina of the spinal cord that receives sensory input from the skin (Zimmermann et al. 2011). It is modulated by receptors coupled to the phosphoinositoside signaling pathway. The use of ratiometric  $\text{Ca}^{2+}$  imaging of dissociated cultured DRG from WT and TRPC5<sup>-/-</sup> mice has demonstrated that TRPM8 and/or other menthol-sensitive channels appear to underpin a much larger component of noxious cold sensing after TRPC5 deletion (Zimmermann et al. 2011).

### 1.3.3 Gating by Temperature and by Voltage in Thermosensitive TRP Channels

As seen above, temperature change in the appropriate range can activate eight heat-activated channels (TRPV1–4, TRPM2–5) and three cold-activated channels (TRPM8, TRPA1, TRPC5) (Clapham 2003; Jordt et al. 2003; Dhaka et al. 2006; Venkatachalam and Montell 2007; Zimmermann et al. 2011). What happens to these channels, however, when temperature changes? A possible answer can be found in the lipids tightly bound to the proteins or surrounding them, thus, these molecules could serve as temperature sensors converting thermal energy into protein conformational changes. However, TRPV1 and TRPM8 channels remain activated by temperature, following perturbations of membrane fluidity by either cholesterol changes (Liu et al. 2003) or by the reconstitution of TRPM8 and TRPV1 into planar lipid bilayer membranes and artificial liposomes, respectively (Zakharian et al. 2010; Cao et al. 2013a).

The key domains or residues for thermosensitivity have not yet been identified and the situation is at present quite unclear. Some studies reveal that the C-terminal of thermo TRPs may be a potential candidate because: (1) swapping the C-terminal domain of TRPV1 with TRPM8 exchanges their temperature sensitivity, channel gating kinetics and PIP<sub>2</sub> modulation (Brauchi et al. 2006, 2007); and (2) sequential deletions of the distal half C-terminal in TRPV1 were shown to shift the temperature threshold to lower temperatures (Vlachova 2003). On the other hand, the pore region has also been postulated as a potential temperature sensor, since it was found that point mutations in the vicinity of the pore region were able to significantly decrease TRPV1 and TRPV3 temperature sensitivity without affecting their activation by voltages (Grandl et al. 2008, 2010). Finally, the TRPV1 N-terminus and *Drosophila* TRPA1 have also been identified as possible thermosensitive structures. Specifically, temperature-dependence has been associated with regions flanking the N-terminal ARD (Cordero-Morales et al. 2011; Yao et al. 2011).

Several TRP channels have been reported to be voltage-sensitive, specifically TRPV1, TRPV3, TRPM8, and TRPM4 (Nilius et al. 2003, 2005b; Voets et al. 2004; Chung et al. 2005). Their voltage sensitivity is quite low in comparison to the classic voltage-gated channels, such as Shaker channels, where movement of S4 across the transmembrane electric field contributes to what would be equivalent to about 13 e<sub>0</sub> gating (Schoppa and Sigworth 1998). What temperature does is shift the voltage activation curve towards more negative voltages according to the heat or cold thermosensitivity of the channels with an unusually large  $Q_{10}$  for a protein. This general allosteric mechanism (similar to the role of intracellular Ca<sup>2+</sup> for BK/Slo1 channels) is also modulated by many intra and extracellular ligands, making TRP channels special sensors and transducer molecules, where many signals can be integrated at the same time, thus behaving like polymodal receptors (Baez-Nieto et al. 2011).



TRPM8, for example, has an apparent gating charge of only 0.6–0.8  $e_0$ . One possible explanation for this could be the scarcity of positive charges in the human TRPM8 S4 domain, having only one arginine residue on the S4 linker (R842) and one lysine on the S4-S5 linker (K856) (Voets 2007). This domain also contains a histidine residue that can be positively charged depending on its pK in the protein milieu. Voets (Voets 2007) designed different charge-neutralizing S4 and S4-S5 linkers for TRPM8, which reduce the channel's gating charge, indicating that this region would be part of the voltage sensor. Similar to TRPM8, TRPV1 presents an apparent gating charge of 0.5–0.7  $e_0$  (Brauchi 2004; Voets et al. 2004, 2007). This channel does not have charged residues in the predicted S4 segment; however it contains an arginine-glutamate pair in the S2 segment (R474 and E478 in rat TRPV1).

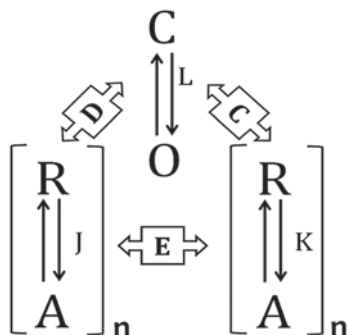
As can be seen, there seems to be no consensus regarding the localization of the temperature and voltage sensors. It is clear, however, that they both reside in different channel molecular structures, because it is possible to eliminate the temperature sensitivity of a channel without altering its voltage sensitivity and vice versa (Boukalova et al. 2010; Yang et al. 2010).

### 1.3.4 Allosteric-Polymodal Gating in ThermoTRP Channels

In the last decade, significant progress has been made in the biophysical and thermodynamic characterization of some thermoTRP channels, especially of TRPV1 and TRPM8 (Brauchi et al. 2004; Voets et al. 2004; Hui et al. 2005; Matta and Ahern 2007). Nevertheless, there is still controversy as to which kinetic model best explains the behavior of these channels. While Voets points out that the two-state model represents a good approximation to describe TRPM8 and TRPV1 temperature and voltage gating (Voets et al. 2004, 2007), analyses of conductance-voltage curves or of single channel data (Fernandez et al. 2011) indicate that this model is an oversimplification. Brauchi made the simplifying assumption that voltage and temperature sensors move in concert, giving rise to a cubic eight-state kinetic model (Horrigan and Aldrich 2002; Brauchi 2004; Latorre et al. 2007a).

The term *allosteric* is quite popular in enzyme kinetics. A typical way of using it in this sense is when a ligand binds to a regulatory site of an enzyme and alters the form of the protein, resulting in changes in enzyme effectiveness. The regulatory site in the protein can be altogether different from the catalytic site, where the chemistry takes place. Thus, to explain TRP channel data (mainly for TRPM8 and TRPV1), an allosteric activation mechanism that involves the following three separate two-state equilibria has been used: voltage sensor activation, temperature sensor activation, channel opening, as well as some interactions among these three processes (Fig. 1.3).

One of the key features that defines the behavior of TRPM8 and TRPV1 channels is that neither temperature nor voltage would be strictly necessary for channel



**Fig. 1.3** Allosteric kinetic model of thermo-RP channel gating. The channel undergoes transitions between closed (*C*) and open (*O*) states with an equilibrium constant *L*, being the voltage and temperature sensors in resting condition. In addition, the sensors undergo transitions between resting (*R*) ((Q9BX84); TRPM7 (Q96QT4); TRPM8 (Q7Z2W7)). Alignments were performed with the cluster method using the UniProt service (ClustalO.) and activated (*A*) states with equilibrium constants *J* and *K* for the voltage and temperature sensors, respectively. The voltage and temperature sensors are coupled to channel gating by allosteric coupling constants *C* and *D*, respectively. The allosteric factor, *E*, establishes the interaction between the voltage and temperature sensors, and *n* represents the number of sensors involved in channel activation. *R*, Resting state we must be careful not to confuse this with the ideal gas constant (*R*)

activation (Brauchi 2004; Latorre et al. 2007a; Matta and Ahern 2007). Thus, at high temperatures, the channel is activated by significant depolarization and when all voltage sensors are resting, the open probability of the channel can be increased by lowering temperature. Furthermore, the allosteric model demands that temperature and voltage sensors be contained in different structures. This has been corroborated by different findings, mainly in TRPM8 and TRPV1 channels, as follows:

1. Specific mutations introduced into the C-terminal of TRPM8 channels abolished temperature sensitivity, yet maintained voltage dependence (Brauchi et al. 2007).
2. Grandl and coworkers (Grandl et al. 2010) found three mutations in TRPV1, localized in the pore region (N628K, N652T and Y653T), which decrease their heat sensitivity as determined by a right shift in the temperature threshold. The double and triple mutants N652T-Y653T and N628K-N652T-Y653T have a stronger phenotype than the single-point mutants, showing greater decreases in their heat sensitivity without affecting their activation by agonists (capsaicin and 2-APB), voltage and acid pH.
3. Analyses of current activation time courses at different temperatures in TRPM8 channels show that they become mono-exponential after brief delays (Latorre et al. 2007a). Such delays are shortened when pre-pulses reach more depolarized values (Cole and Moore 1960), indicating that the channels transit between several voltage-dependent closed states. Similarly, large hyperpolarizing voltages would populate the closed states further removed from open states. These

changes in delays do not show temperature dependence, suggesting that temperature dependence of voltage-dependent transitions is weak.

- Analyses of single channel data indicate the existence of a minimum of five closed states and two open states for TRPM8 (Fernandez et al. 2011) and at least three closed and three open states for TRPV1, whose duration is practically temperature-independent (Liu et al. 2003).

All these results strongly suggest that voltage and temperature sensors are different entities. Allosteric models have been successful in explaining the workings of other polymodal ion channels, such as BK potassium channels (Horrigan and Aldrich 2002; Orío and Latorre 2005) and can be useful to describe the complex regulation of thermo TRP channels, such as TRPV1 and TRPM8, and to understand their contribution to heat and cold sensitivity.

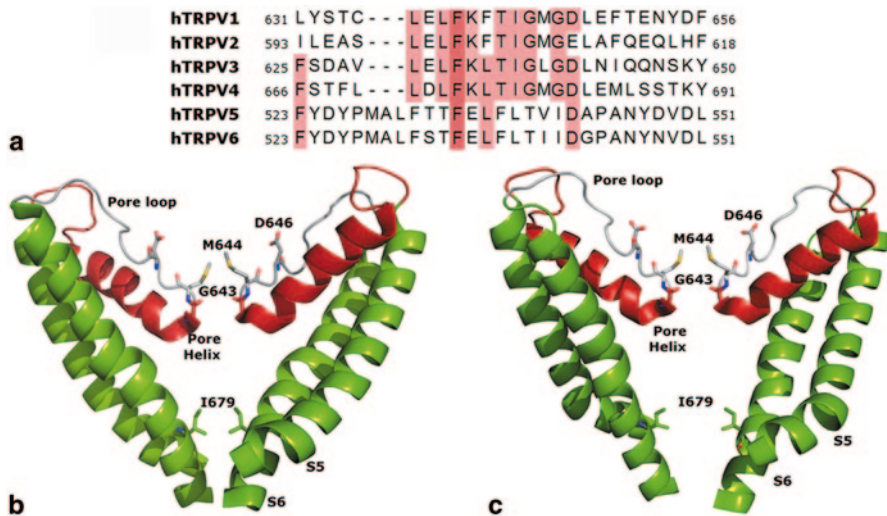
### ***1.3.5 Pore and Selectivity of Thermosensitive TRP Channels***

TRP channels are non-selective cation channels, permeable to both monovalent and divalent cations. However, the degree of specificity of permeation to either monovalent or divalent cations depends on the type of TRP channel. In some cases, it seems that the permeation properties of a channel may be associated with its complex polymodal gating, as it has recently been shown by Cao et al. (Cao et al. 2013b).

#### **1.3.5.1 The TRPV Subfamily**

While TRPV1–4 channels are mostly involved in nociception and thermal reception, TRPV5/6 channels are mostly involved in  $\text{Ca}^{2+}$  reabsorption in the kidney (Owsianik et al. 2006). Thus, TRPV5/6 channels are highly permeable to  $\text{Ca}^{2+}$ , whereas TRPV1–4 channels are mildly  $\text{Ca}^{2+}$  permeable. They also exhibit rectification and modulation by divalent cations and protons. In contrast to TRPV 5/6 channels that show strong inward rectification in the absence of divalent cations (Voets et al. 2003), TRPV1 exhibits outward rectification (Premkumar et al. 2002; Voets et al. 2003) and extracellular divalents can potentiate the TRPV1 current, promoting changes in the extracellular region (Yang et al. 2014).

Alignments of TRPV channels show that there are conserved residues (L, F, I; see Fig. 1.4a) in the putative pore region among all these subfamily members, and also that the mutations of aspartates have profound effects on the selectivity and blocking properties of TRPV4–6 channels (Nilius et al. 2001; Voets et al. 2002, 2003). Accordingly, it has been proposed that the aspartates in these channels would form a tetrameric ring with two binding sites for  $\text{Ca}^{2+}$  (Nilius et al. 2001; Voets and Nilius 2003). Thus, D646 has been identified to be relevant for  $\text{Ca}^{2+}$  selectivity and blocking by Ruthenium Red in TRPV1 (Garcia-Martinez et al. 2000). Structure-function studies in TRPV1 channels have revealed that the S5-S6 loop is related to the permeation pathway (Lopez et al. 1994). Recently, electron cryo-microscopy



**Fig. 1.4** The putative pore structure in TRPV channels subfamily. **a** Sequence alignment of the pore region of human TRPV channels subfamily. The conserved amino acids that in all or mostly members of subfamily are indicated in *red*. Sequences used in the alignments were extracted from UniProt database: TRPV1 (Q8NER1); TRPV2 (Q9Y5S1); TRPV3 (Q8NET8); TRPV4 (Q9HBA0); TRPV5 (Q9NQA5); TRPV6 (Q9H1D0). Alignments were performed by cluster method using UniProt service (ClustalO). **b** Molecular structure of the rat TRPV1 pore region in closed (PDB 3J5P) and **c** open states (PDB 3J5R), showing pore loop (Trevisani et al.) and pore helix domain (Zaritsky et al.), selectivity filter (GMDG), and lower gate (I679). Transmembrane domain 5 (S5) and 6 (S6) of adjacent subunits are represented in *green*

has been applied to the TRPV1 channel with an average resolution of 3.4 Å, which was revealed to have a wide outer pore followed by a short-narrow selectivity filter (G643-M644-G645-D646) and a lower gate (I679) (Liao et al. 2013) (Fig. 1.4b).

As stated before, temperature gating in TRPV1 (and TRPM8) is allosterically linked to gating by chemicals and by voltage (Brauchi et al. 2006; Latorre et al. 2007a). It has also been shown that thermal or chemical stimulations of some TRPV channels have profound effects on their pore properties and structure, especially the outer pore region (Bautista and Julius 2008). Recent studies have shown that single pore residues in an extension of 50 amino acids in the outer mouth of the pore in both TRPV1/3 channels exhibit temperature-dependent conformations (Kim et al. 2013). However, some authors have reported significant conformational changes of the pore turret in TRPV1 channels due to temperature (Yang et al. 2010), whereas others implicate the inner pore region as one of the critical domains determining TRPV1 channel gating, also modulating capsaicin and thermal sensitivity (Susankova et al. 2007; Myers et al. 2008). Differences in the pore region may account for the differences observed between species for TRPV1 heat activation (Papakosta et al. 2011).

Finally, modulation by protons in TRPV5 has been shown to alter pore dimensions by cysteine scanning mutagenesis. Extra or intracellular acidification has been seen to facilitate pore collapse in TRPV5 through pore helix rotations (Yeh



tusa et al. 2012). The site is flanked by two cysteines (C929 and C940), which are essential for channel function (Dragoni et al. 2006) (Fig. 1.5).

### 1.3.5.3 The TRPA Subfamily (TRPA1)

While TRPM8 is activated by moderate cold (Temp < approx. 25 °C), TRPA1 is activated by noxious cold (<17 °C) (Story et al. 2003). These channels are quite permeable to Ca<sup>2+</sup>, being the third among TRP channels (TRPV5/6 channels being the most permeable to Ca<sup>2+</sup>). D918 and E923 seem to be critical for Ca<sup>2+</sup> permeation (Karashima et al. 2010). As it has been reported for TRPV channels, there have been studies linking the pore and gating of these channels to temperature (Benedikt et al. 2009; Banke et al. 2010; Wang et al. 2013). Using organic cations of different sizes, it has been shown that activation of TRPA1 by cold involves reversible pore dilation, yielding pore diameters of about 11 Å (Chen et al. 2009; Banke et al. 2010; Alpizar et al., 2013).

### 1.3.5.4 The TRPC5 Channel

TRPC5 modulation by temperature was unknown until 2011, when it was established as a cold-activated channel in the peripheral nervous system (Zimmermann et al. 2011). Homotetrameric TRPC5 channels show an unusual biphasic rectification property that might be explained by voltage-dependent blocking by intracellular Mg<sup>2+</sup> (Obukhov and Nowycky 2005). Unlike other TRP channels, the region between S5 and S6 in TRPC channels is not conserved. Negative residues inside and nearly outside the S5-S6 loop, like E559/E570, might be related to the selectivity properties of the TRPC5 channel, suggesting that the pore region in this subfamily is also related to the S5-S6 loop and helices, as in the rest of the TRP channel families (Gutman et al. 2003).

## 1.3.6 *Gating Modifiers in the Context of Molecular Models of Thermo TRP Channels*

*TRPV* channels are voltage-activated channels, whose voltage thresholds for channel opening can be allosterically modified by heat (temp. >42 °C), capsaicin and protons in an allosteric polymodal way (Latorre et al. 2010; Baez-Nieto et al. 2011). It has been known that TRPV1 channels are activated by pH < 6 (Caterina et al. 1997). Lowering the pH was seen to diminish the threshold for activation by capsaicin and related compounds as well as heat. Proton binding residues in TRPV1 were characterized as E600 and 648, and they were shown to be relevant for TRPV1 channel modulation by other extracellular cations, as well as for the modulation of sensitivity to chemical agonists (Jordt et al. 2000; Ahern et al. 2005; Ryu et al. 2007)



*Capsaicin* and its natural and artificial derivatives (capsacinoids) activate the TRPV1 channel and they seem to do so from within the intracellular moieties of the channel. Sensitivity to capsaicin and its endogenous analog, anandamide, was shown to be closely related to transmembrane segment 3 (S3 or TM3). Based on site-directed mutagenesis of TRPV1, Jordt and Julius (Jordt and Julius 2002) proposed that tyrosine at position 511 (Y511) and its aromatic nature, located in or nearby the intracellular TM3 loop, is critical for TRPV1 capsaicin sensitivity. Residues close to Y511, like S512 or R491, have been observed to have an important influence on capsaicin activity as well. These results led the authors to propose that all such residues constitute a “pocket” for vanilloid-like agonist molecules.

The *N-terminal* cytosolic segment of TRPV channels contains ankyrin repeat domains (Stewart et al. 2010), whose sequences are well conserved. The ARDs have been shown to be important for channel gating modulation and sensitization through ATP binding and  $\text{Ca}^{2+}$ /calmodulin interaction (TRPV1–4, but not TRPV2/5–6) (Rosenbaum et al. 2004; Lishko et al. 2007; Phelps et al. 2010). In TRPV1/3 channels,  $\text{Ca}^{2+}$ -calmodulin binding to ARDs in addition to a C-terminal binding site promote desensitization, though it potentiates TRPV4 channels (Lishko et al. 2007; Phelps et al. 2010; Lau et al. 2012). Sensitization in TRPV3 takes place when  $\text{Ca}^{2+}$  is released from the inhibiting binding sites located in the ARD calmodulin interaction site (residues 108–130), as well as the release of  $\text{Ca}^{2+}$  from the pore to the extracellular side (Asp 641 inside the pore) (Xiao et al. 2008). Regarding TRPV4, it has been suggested that potentiation may arise from interactions between the N-terminus and C-terminus  $\text{Ca}^{2+}$ -calmodulin binding-sites (Strotmann et al. 2003, 2010). All these results indicate that despite the sequence conservation among thermosensitive TRPV1–4 channels, they are differentially modulated by the same ligands and sites, suggesting different 3D structures that might be explained by distinct fits of the ARDs inside the cytoplasmic side of the channel (Huynh et al. 2014).

The *C-terminus* (the proximal region to S6) contains a very well conserved sequence among all TRP channels of 25 residues, named TRP domain where the *TRP-Box* (IWKLQR) (Ramsey et al. 2006). The following coiled-coil region (E684 to R721), also comprising part of the rat TRP domain, is involved in TRPV1 tetramerization (Vlachova et al. 2003; Garcia-Sanz et al. 2004). Similar residues have been identified for mouse TRPV1 (752–772), receiving the name *tetrameric assembly domain* (Wirkner et al. 2005) (Zhang et al. 2011). Furthermore, in TRPV1 it has been established that this region might participate in calmodulin-binding (E767 to T801), phosphorylation and thermal sensitivity (Numazaki et al. 2003; Brauchi et al. 2006; Huang et al. 2006).

$\text{Ca}^{2+}$ -calmodulin binding has been implicated in TRPV1 desensitization (Numazaki et al. 2003). However, the C-terminus calmodulin-binding site (named CT, from W787 to L796) has distinct effects on desensitization compared to the N-terminus ARD calmodulin-binding site (Lau et al. 2012). Interestingly, the CT calmodulin-binding site does not contain a classic calmodulin binding sequence motif. The crystallized 3D structure of the TRPV1-CT segment shows that it is quite similar to the canonical  $\text{Ca}^{2+}$ -calmodulin peptide complexes, having two lobes of  $\text{Ca}^{2+}$ -calmodulin wrapped around the TRPV1-CT segment (Lau et al. 2012). Unlike

TRPV1, the calmodulin interaction site in TRPV2 has been characterized as a classic calmodulin binding sequence (Holakovska et al. 2011) and  $\text{Ca}^{2+}$ -desensitization is through  $\text{PIP}_2$  hydrolysis (Mercado et al. 2010). Furthermore, differential thermal sensitivity shorting this region in TRPV1 channels was reported by Vlachova et al. (Vlachova et al. 2003).

Bearing all of this in mind, Brauchi et al. (2006) switched the C-terminal tail between hot-sensing TRPV1 and cold-sensing TRPM8. Switching the channel's tail resulted in a chimeric channel, whose phenotype was determined by the C-terminal tail, not only for thermal sensing, but also for some gating properties such as modulation of channel function by  $\text{PIP}_2$  binding. Recently, it has been shown that  $\text{PIP}_2$  activates the channel by directly interacts with TRPV1, and also that the  $\text{PIP}_2$  binding-site is located proximal to the TRPV1 channel C-terminus (Ufret-Vincenty et al. 2011). The precise interaction location of  $\text{PIP}_2$  with the C-terminus has been reported to take place at residues K688-K718 and L777-S820, constituting two independent sites that overlap with the CT segment. In addition, a third binding site for  $\text{PIP}_2$  has also been reported in the TRPV1 channel N-terminus (F189-V221) (Grycova et al. 2012).

**TRPM channels** are also voltage-activated, just like TRPV channels. In contrast with TRPV1–4 channels, the TRPM8 channel opening voltage threshold is activated by cold (temp.  $<25^\circ\text{C}$ ) as well as by menthol and derivatives. High-throughput screening of random mutagenesis in mouse TRPM8 has identified a tyrosine residue in S2 (Y745), which is critical for *menthol* sensitivity (Bandell et al. 2006). It has been suggested that there should be many other residues in the channel contributing to the menthol-binding site, as some TRPM8 antagonists that supposedly interact with the menthol-binding site are still active in Y745 TRPM8 mutated channels (Malkia et al. 2009). Mutations in S4 (R842H) can also change menthol sensitivity, suggesting that its site of action could be placed in the cleft between S2 and S4 (Voets et al. 2007). Subthreshold concentrations of menthol move TRPM8 activation cold temperature threshold to warmer temperatures. However, repeated menthol stimulation rapidly desensitizes TRPM8 in a  $\text{Ca}^{2+}$ -dependent manner (McKemy et al. 2002; Reid 2002).

TRPM8 is also activated by the so-called “super-cooling agent” *icilin* (as it is 200 times more potent than menthol) in the presence of intracellular  $\text{Ca}^{2+}$  (Chuang et al. 2004). The binding site for icilin seems to be located in residues along S3 (N799, D802 and G805). This finding implies that icilin and menthol-binding sites are different and that they would also probably differ in terms of their channel activation mechanisms (Chuang et al. 2004). In line with this view, it has been shown that icilin and cold, but not menthol sensitivity, is modulated by intracellular pH and viceversa (Andersson et al. 2004).

As discussed above, the intracellular *N-terminus* of TRPM channels has a large conserved region among channel subfamily members (~700 residues), and are characterized by having 4 subdomains labeled TRPM homology regions or *MHRs* involved in trafficking (TRPM2/4/8) (Clapham 2003; (Perraud et al. 2003; Launay et al. 2004; Phelps and Gaudet 2007).



In TRPM2/4/5 channels, activity critically depends on the level of intracellular  $\text{Ca}^{2+}$  and has been implicated in channel gating and desensitization (McHugh et al. 2003; Tong et al. 2006); Vennekens and Nilius 2007 (Nilius et al. 2004a). In TRPM2, intracellular  $\text{Ca}^{2+}$  activates the channel by binding to calmodulin, presumably located at this IQ sequence motif in the N-terminus of the channel (residues 406–416) (McHugh et al. 2003; Tong et al. 2006).  $\text{Ca}^{2+}$ -sensitivity is also determined by the binding of four  $\text{Ca}^{2+}$  ions, which are cooperatively bound to the inner mouth of the channel in the open state (Csanady and Toroicsik 2009). In TRPM4, ATP binding to the N-terminus and to the S3-S4 intracellular loop appears to be needed for  $\text{Ca}^{2+}$  channel sensitivity. Calmodulin binding sites in the N-terminus, however, appear to be irrelevant for TRPM4 function (Nilius et al. 2004b).

Like TRPV channels, the *C-terminus* contains the *TRP domain* and box that is well conserved among all TRPM channels (McKemy et al. 2002). The C-terminus distal region contains a coiled-coil structure involved in multimerization, trafficking and activation gating of TRPM8 (residues 1064–1104), which is conserved among TRPM channels (Erler et al. 2006; Mei et al. 2006; Tsuruda et al. 2006; Phelps and Gaudet 2007).

Regulation of thermoTRPM channels by PIP<sub>2</sub> includes activation or potentiation of TRPM4/5/8 (Guinamard et al. 2011; Prescott and Julius 2003; Vennekens and Nilius 2007). In TRPM8, some key positively charged residues present in the TRP box (K995, K998 and R1008) were found to be crucial in PIP<sub>2</sub> binding and modulation (Rohacs et al. 2005).  $\text{Ca}^{2+}$ -dependent desensitization of TRPM8 seems to occur through hydrolysis of PIP<sub>2</sub> (Liu and Qin 2005; Rohacs et al. 2005). Y1005 and L1009 residues have been implicated in the modulation of menthol sensitivity, but they do not seem to be involved in the initial event needed for menthol activation (Bandell et al. 2006). The results obtained with splice variants of the TRPM8 channel C-terminus (named  $\alpha$  and  $\beta$ ) suggest that these short isoforms seem to mimic heat, hence inhibiting channel activation. Therefore, all of these results are consistent with the hypothesis that the C-terminus in TRPM8 channels is essential for their cold sensing ability, channel activation and modulation by PIP<sub>2</sub> (Brauchi et al. 2006; Bidaux et al. 2012; Fernandez et al. 2012).

**TRPA1 channels**, as we discussed above, are quite indiscriminating when it comes to their gating modulation. In addition to the agents that commonly modulate most of the TRP channels described in this review (such as  $\text{Ca}^{2+}$ , phosphorylation/dephosphorylation and PIP<sub>2</sub>), TRPA1 channels are also modulated by a plethora of different compounds that have pharmacological and physiological significance. These natural or artificial chemicals can produce nucleophilic covalent modifications (reviewed in (Nilius et al. 2012)) (i.e., electrophilic compounds such as isothiocyanates, cinnamaldehyde, allicin, acrolein, O<sub>2</sub>, CO<sub>2</sub>, reactive oxygen species (ROS) and reactive nitrogen species (Watanabe et al. 2002a)). Other types of interactions like non-electrophilic compounds (menthol, icilin, caffeine) are unlikely to modify channel gating through covalent interaction, exhibiting bimodal action (bell-shaped dose response curves) (Nilius et al. 2012). Thus, activation by cold in TRPA1 is potentiated in the presence of many of the chemical agonists mentioned above (Fajardo et al. 2008).

The *N-terminus* is characterized by having a long stretch of 18 ARDs, which are important for protein-protein interaction and channel trafficking. Intracellular  $\text{Ca}^{2+}$  can bind to the N-terminus and change its structure (Nilius et al. 2012). However, the deletion of the putative EF-hands that might act as N-terminal  $\text{Ca}^{2+}$ -binding sites only have effects on TRPA1 membrane trafficking (just mild effects on the modulation of TRPA1 by intracellular  $\text{Ca}^{2+}$ ) (Nilius et al. 2011). There are cysteine residues in the N-terminus that could be quite important for TRPA1 interaction through disulfide bridges, because they are located at the interface between TRPA1 subunits, hence being able to interfere with gating mechanisms. In addition, Sura et al. (2012) reported that site-directed mutagenesis in the *C-terminus*, which contains a stretch of several negative amino acids arranged in a manner similar to BK  $\text{Ca}^{2+}$ , impairs TRPA1 modulation by means of intracellular  $\text{Ca}^{2+}$  (residues E1077, D1080, D1081 and D1082) (Sura et al. 2012). Other residues like Lys 1048, Lys 1052, Lys 1092 and Arg 1099 have also been observed to modify a significant amount of TRPA1 chemical and voltage-dependent activation gating. These basic residues in the C-terminus might also explain the diversity of electrophilic compounds that can affect TRPA1 channel gating (Samad et al. 2011).

**TRPC5 channels** are present in mouse and human sensory neurons of dorsoal root ganglia, a substantial number of peripheral nerves, and in dorsal lamina of the spinal cord. These channels are non-selective cation channels permeable to  $\text{Ca}^{2+}$  and modulated by a large variety of signals, as it is usually reported for TRP channels (Zeng et al. 2004): intracellular  $\text{Ca}^{2+}$ , ATP, protons,  $\text{PIP}_2$  and phosphorylation/dephosphorylation (Semtner et al. 2007; Dattilo et al. 2008). Some of these mechanisms are complex and seem to require different sites of action. In particular, it has been reported that intracellular  $\text{Ca}^{2+}$  usually potentiates agonist-activated TRPC5 channels (Blair et al. 2009). The *C-terminus* seems to contain  $\text{Ca}^{2+}$ -binding protein interaction sites, through which intracellular  $\text{Ca}^{2+}$  may exert its effects (calmodulin,  $\text{Ca}^{2+}$  binding protein 1, NCS-1) (Ordaz et al. 2005; Shimizu et al. 2006; Beech 2007).

In addition, these channels are activated by hypo-osmotic and pressure related changes in the membrane, revealing their role in mechanosensation (Gomis et al. 2008). Some of these changes are potentiated upon G-protein coupled receptor (GPCR) activation (Jemal et al. 2013). G proteins released from the GPCR increase channel responses to cold (Zimmermann et al. 2011) and might mediate PKA action in TRPC5 (Sung et al. 2011). Sung *et al* identified serines located in the C-terminus (S794 and S796) as a PKA phosphorylation site. TRPC5 channels are also strongly modulated by oxidizing agents that act through nitrosylation, like nitric oxide (NO). They can activate the channel by reacting with cysteine residues located at 553–558 in the S5-S6 linker, which are accessible from intracellular media (Yoshida et al. 2006).

### 1.3.7 Evolution of Thermosensitive TRP Channels

Since evolution involves a selection of the most adapted individuals to the environment in which they live, TRP channels are critical for such adaptation because of their nature as sensing and integrating molecules. Studies on TRP channels from an evolutionary perspective are very relevant to the field, as information about sites for particular chemical or physical stimuli can be obtained by comparing ortholog and paralog genes in different species that appeared on Earth at different times. They may also reveal critical interaction sites necessary for TRP channel function and interaction with other proteins (Sardar et al. 2012). The phylogenetic tree for the seven subfamilies of TRP channels (mostly including human TRP channels) is shown in Fig. 1.2 above.

TRP channels have only been described in protozoa (*Leishmania*), fungi and animalia thus far (Prole and Taylor 2011; Wolstenholme et al. 2011). Though TRP channels are not endogenously expressed in prokaryotes (Li et al. 2011), bacteria still have ways for osmo and mechano-transduction, which are not as elaborated as in eukaryotes, where channel sensing molecules functioning as precursors of TRP channels are involved (Jin et al. 2011). TRPN channels are only present in invertebrates and some fish, and absent in most vertebrates, which only have six subfamilies. TRPN channels are named as such because of the “No-mechano-potential-C” channels from *C. Elegans*. Although these channels (TRPN) were lost in vertebrates during evolution, the TRP channels almost doubled in numbers when implicated in  $\text{Ca}^{2+}$  and  $\text{Mg}^{2+}$  homeostasis as well as in thermosensation (Nilius and Owsianik 2011). In this regard, sponges, which are one of the most ancient metazoans with active responses to stimuli, though lacking nervous systems, contain a cilia-sensing system with TRP channels (Ludeman et al. 2014).

Animal adaptation to temperature changes occurs by sensing both body and environmental temperature. Toward this aim, thermal stimuli are monitored by thermal receptors expressing thermoTRP channels. Hence, changes in thermoTRP channels and thermosensitivity are crucial in understanding animal adaptation to different environments with different temperatures. Most of these genes emerged from common teleost fish ancestors through gene duplication (Saito and Shingai 2006). Regarding TRPVs, thermo-TRPVs (TRPV1–4) and nonthermo-TRPVs (TRPV5–6) diverged early in evolution (Fig. 1.2). ThermoTRPVs split first into the TRPV4 and TRPV1–3 branches, and later divided into the TRPV1–2 branch and TRPV3 channels (Saito et al. 2011). TRPV3–4 channels diverged earlier than the separation of the TRPV1–2 branch. TRPV1–2 channels are activated by noxious heat (temp.  $>40^\circ\text{C}$ ), whereas, TRPV3–4 are activated by warm temperatures ( $T > 35^\circ\text{C}$ ). Because of that, they have been implicated in temperature regulation in homeothermic vertebrates. Consistent with this idea, TRPV3-null mice lack adequate control of their body and environmental temperatures (Moqrich et al. 2005). A TRPV3 ortholog of the mammalian TRPV3 channel found in western claw frogs has been seen to significantly diverge from mammalian TRPV3 channels in their C-terminus and N-terminus. In addition, they lack responses to several agents known to modulate mammalian TRPV3. All these results are consistent with previous findings that

suggested critical roles of the N-terminus and C-terminus in TRPV channels in heat and chemical modulation (Saito et al. 2011; Brauchi et al. 2004).

These results are quite interesting, since they show that the range of temperature sensitivity in thermoTRP channels can be changed and adapted during evolution, hence being flexible. Unlike mammals, which are homeothermic vertebrates with body temperatures around 35–37°C, western clawed frogs are ectothermic vertebrates adapted to lower temperatures. Western clawed frogs live in an environment with temperatures within the range of 22–28°C. Just like mammals, in which TRPV3 senses adequate temperatures in the body surface, TRPV3 in the western clawed frog is also crucial in perceiving skin temperature (Mandadi et al. 2009; Saito et al. 2011). In this sense, TRPV3 channels in western clawed frogs lack the ability to be modulated by heat and cold, and tend to mainly detect noxious cold, as it is of more advantage for the survival of the frog-environment relationship. The results are also consistent with the idea of modular “hot spots” in the functional structure of TRP channels, related to their different sensing abilities (Latorre et al. 2007a).

Opposite thermosensitivity among other ortholog thermoTRP channels besides TRPV3 has been reported in TRPA1 channels. While such thermosensitivity is activated by warm or hot temperatures in reptiles, frogs and insects, it is activated by cold in mice and it is not activated by temperature in zebrafish or humans (Viswanath et al. 2003; Kwan et al. 2006; Karashima et al. 2009; Gracheva et al. 2010; Saito et al. 2012); Chen et al. 2013). TRPA1 channels are quite ancient among animals and are interesting in that their thermal sensitivity was shifted from hot to cold during the course of evolution. Invertebrates like *Drosophila melanogaster* and *C. elegans* show that TRPA1 is sensitive to heat and nociceptive chemical stimulation (Wang et al. 2009; Kang et al. 2012). Except for zebrafish, where TRPA1 is insensitive to temperature, early vertebrates like amphibians, reptiles and lizards possess TRPA1 channels sensitive to heat with variable thresholds. In all cases, nociception by chemicals was maintained through time (Prober et al. 2008). Swapping the N-terminus of snake and mammal TRPA1 channels seemed to have changed their temperature dependence (Cordero-Morales et al. 2011). This suggests that the module related to TRPA1 channel thermosensitivity could be placed in the N-terminus, whilst nociception by chemicals would be due to binding to different regions of TRPA1 channels.

TRPA1 and TRPV1 are co-expressed in the same cells and might form heteromers as well (Salas et al. 2009; Staruschenko et al. 2010; Akopian 2011). TRPV1 came from the TRPV1/2 branch separation later in evolution after TRPA1 (Saito and Shingai 2006). Once TRPV1 channels appeared, they maintained their gating characteristics by heat as well as their temperature thresholds through the span of evolution. It has been hypothesized that the appearance of TRPV1 channels in evolution after TRPA1 channels as efficient sensors for noxious heat and nociception favored the transition of TRPA1 channels as receptors of noxious cold, while maintaining their nociception abilities. Hence, both channels are frequently expressed together in neurons related to nociception (Saito et al. 2012).

## 1.4 Concluding Remarks

In this chapter we have reviewed the structural, biophysical and evolutionary details of thermoTRP channels, relevant for sensory transduction. Regarding their biophysics and physiology, their allosteric polymodal nature confers them, among other proteins, an enormous advantage for sensing and transducing signals. The wide repertoire of chemical and physical agents that are able to gate these channels provides sophisticated fine-tuning for cell and organism living conditions. Their non-selective cation permeation properties and their mild voltage gating, which are strongly modulated by chemical and physical agents that can interact allosterically, make them unique machineries for cell homeostasis. Regarding their structure, their organization as tetramers with interrelated sensing modules has been revealed through structure-function studies. More recently, 3D models have shown that they have large cytoplasmic interacting regions, making them suitable for monitoring intracellular changes interacting with many ligands and proteins. All these features are reflected in evolution as they appear in molecules within more complex organisms. In line with the structural and biophysical details known to date, analyses of their properties in different organisms have revealed that they are one of the key factors involved in adapting to distinct thermal conditions. As such, these unique properties are conferred to cells and organisms expressing these molecules, indicating a clear advantage that makes them suitable for building complex interactions and networks. Network analysis of these channels in physiology and diseases has just begun (Chun et al. 2013) and have contributed in shedding light on our understanding as to how these channels play important roles in life within the context of systems biology.

**Acknowledgements** This work was supported by FONDECYT Grants 1110430 and 1150273 (to Latorre R), 1120802 (to González C); ANILLO Grant ACT1104 (to Gonzalez C). CSIC p944, the CSIC International Cooperation Program, U de la R-CINV-UV and the Chilean Conicyt International Cooperation Program (to Ferreira G and González C). The *Centro Interdisciplinario de Neurociencia de Valparaíso* is a Millennium Institute supported by the Millennium Scientific Initiative of the Chilean Ministry of Economy, Public Works, and Tourism.

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## Chapter 2

# Pharmacology of TRP Channels

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**Abstract** TRP channels are a family of ion channels involved in a plethora of physiological sensory processes. Since their discovery they have attracted the attention of academic and non-academic laboratories with the aim of developing modulators that could be used as pharmacological tools for unveiling their physiological and pathological activities, and as therapeutic compounds for intervening in TRP dysfunction. Intriguingly, TRP pharmacology shows dispersed progress, with vast pharmacology developed for some members of the so-called thermoTRP channel subfamily (TRPV1, TRPV3, TRPM8 and TRPA1), and very little, for all other TRP channels. Pharmacologically, the most investigated TRP channel is undoubtedly TRPV1 for which a large number of agonists and antagonists with *in vitro* and *in vivo* activities have been characterized. Recent interest has grown for TRPV3, TRPM8 and TRPA1 because of their implication in several human pathologies and disorders. Similarly, the TRPM3 channel is emerging as important targets for pain transduction. With the development of novel screening methods, the focus is slowly changing to other TRP members for whom we do not have appropriate agonists or antagonists. These include the TRPC family, which has limited our understanding of their role in pathological processes and whether pharmacological intervention in these channels will have a therapeutic benefit. A bright future is anticipated for TRP pharmacology, with the discovery of selective and potent modulators for this important family of sensory channels.

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© Springer International Publishing Switzerland 2015  
R. Madrid, J. Bacigalupo (eds.), *TRP Channels in Sensory Transduction*,  
DOI 10.1007/978-3-319-18705-1\_2

**Keywords** TRP channels · Agonists · Antagonists · Competitive · Non-competitive · Uncompetitive · Pain · Therapeutic index · Pathology

## 2.1 Introduction

TRP channels are a superfamily of ion channels that includes seven subfamilies, namely TRPC, TRPV, TRPP, TRPM, TRPA, TRPML, and TRPN. These channels perform a wide diversity of physiological functions and are present in many tissues, and almost all cell types. Most TRP channels are non-selective cation channels with low voltage dependence. TRP channels use a wide variety of activation and regulatory mechanisms and carry out functions as diverse as thermosensation, phototransduction, pheromone reception, magnesium homeostasis, and vascular tone regulation (Montell 1999) (see Chap. 4 by Bacigalupo et al. in this Book). Thus, these channels are considered molecular gateways in sensory and regulatory systems.

Structurally, TRP channels are tetrameric assemblies of basic subunits organized around a central aqueous pore. Akin to voltage-gated  $K^+$  channels, each subunit is composed of a transmembrane region containing 6 transmembrane segments. The recent structural model derived from cryo-electron microscopic images has clearly shown this molecular analogy (Liao et al. 2013). All TRP channels display this core transmembrane region, and differ in the cytosolic N- and C-termini domains, which are involved in channel gating and mediating intracellular signaling. Indeed, most of TRP channels, if not all, are part of protein complexes known as signalplexes (Devesa et al. 2011; Fernandez-Carvajal et al. 2011; Ferrer-Montiel et al. 2012).

Some TRP channels have been involved in the pathophysiology of human diseases. This pathological contribution could be the result of channel mutations, giving rise to channelopathies (Devesa et al. 2011; Fernandez-Carvajal et al. 2011; Ferrer-Montiel et al. 2012), or the change in channel function due to alteration of the protein function and/or expression (Devesa et al. 2011; Fernandez-Carvajal et al. 2011; Ferrer-Montiel et al. 2012). The pivotal involvement in the etiology of pathological conditions has signaled members of this large channel family as druggable targets for therapeutic intervention, which has driven discovery programs in academic and non-academic institutions. This concerted effort has notably expanded the pharmacology of TRP channels, although, unfortunately, for a limited number of TRP members. For instance, large families of modulators have been obtained for TRPV1, TRPV3, TRPM8 and TRPA1, while the pharmacology of other TRP channels is still in its infancy. A plausible reason for the pharmacological progress in these channels is the availability of natural ligands present in food spices. Nonetheless, the development of combinatorial chemistry and the large diversity of vegetal and marine extracts, along with the development of high throughput electrophysiological assays for ion channels will expand the pharmacology of TRP channels to the entire family. Here, we briefly expose the pharmacological data for the most studied TRP channels, namely TRPV1, TRPM8 and TRPA1, and include the data accrued for TRPV2, TRPV4, TRPM3 and the TRPC5, most of them with an

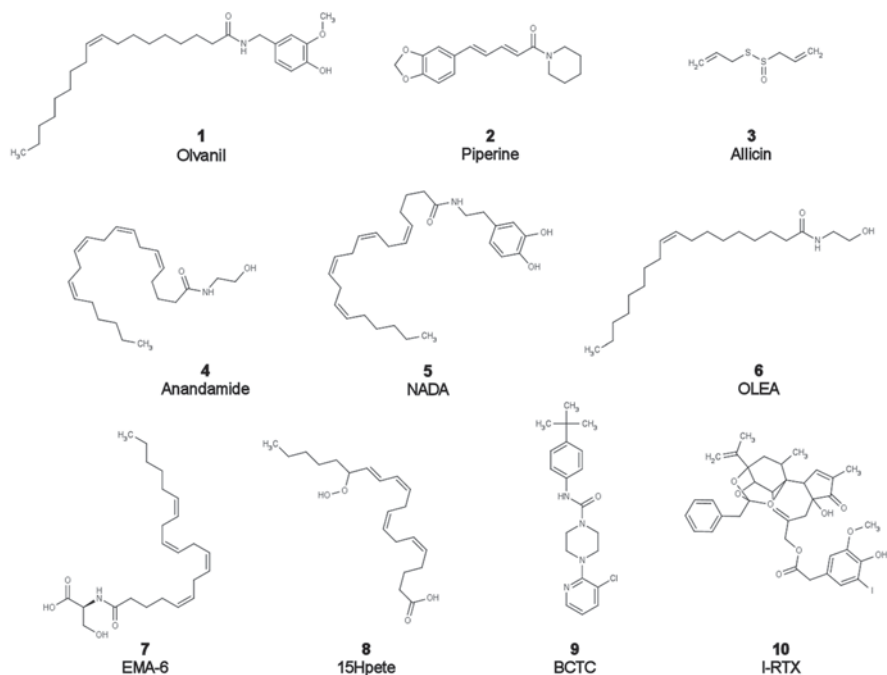
important role in sensory transduction. We aim to illustrate the differential pharmacological progress in this exciting field and evidence a drift towards enhancing the pharmacology of other members, if not all, of this pivotal channel family.

## 2.2 TRPV1

TRP Vanilloid 1, TRPV1, a non-selective  $\text{Ca}^{2+}$  channel is a TRP channel activated by noxious temperatures ( $43^\circ\text{C}$ ) acidic pH and vanilloid compounds, whose channel activity is highly potentiated by proalgesic mediators in response to inflammation, tissue injury and ischemia (Huang et al. 2006; Ueda et al. 2008). In addition, TRPV1 expression is markedly up-regulated under acute inflammatory conditions (Camprubi-Robles et al. 2009; Morenilla-Palao et al. 2004; Van Buren et al. 2005), and in human chronic pain states (Broad et al. 2008; Szallasi and Blumberg 2007). Consistent with a role in pain signaling, TRPV1 is highly expressed in C-type, peptidergic nociceptors in the peripheral nervous system. Thus, TRPV1 is considered a gateway for pain transduction, and a pivotal target for drug intervention in pain syndromes. In addition, due to a widespread tissue distribution of this TRP channel, it may be involved in the etiology of other human pathologies or disorders (Avelino et al. 2002; Inoue et al. 2002).

TRPV1 sensitization by inflammatory conditions is produced through two distinct, but complementary mechanisms, namely: (i) covalent modification of the channel by protein kinase A (PKA) and/or protein kinase C (PKC) phosphorylation (Bhave et al. 2003; Tominaga et al. 2001; Varga et al. 2006; Vellani et al. 2001); and, (ii) rapid recruitment of a vesicular population of TRPV1 channels to the neuronal surface through a  $\text{Ca}^{2+}$ -dependent, SNARE-mediated exocytosis mechanism in response to pro-algesic agents (Camprubi-Robles et al. 2009; Zhang et al. 2005).

Pharmacologically, TRPV1 is primarily activated by a diverse collection of chemical ligands known as vanilloids (Caterina et al. 2000; Khairatkar-Joshi and Szallasi 2009) (Fig. 2.1). The most known agonist of TRPV1 is capsaicin, the pungent compound of chili peppers. Resiniferatoxin (RTX), a vanilloid from *Euphorbia resinifera*, is also a potent agonist of the receptor. Furthermore, TRPV1 may also be activated by non-vanilloid compounds, such as allicin, piperine, camphor, olvanil, 2-aminoethoxydiphenylborate (2-APB), and tarantula venom peptide toxins (Bohlen et al. 2010) (Table 2.1). In addition, there is a family of endogenous compounds, referred to as endovanilloids, that also act as agonists of TRPV1 (Van Der Stelt and Di 2004). These compounds may be divided into conjugates of biogenic amines [e.g., N-arachidonoylthanolamine (AEA, anandamide), N-arachidonoyldopamine (NADA), N-oleoylthanolamine (OLEA), N-arachidonoylserine, and various N-acyltaurines and N-acylsalsolinols (Appendino et al. 2008), and oxygenated eicosatetraenoic acids like the lipoxygenase products 5-, 12-, and 15-hydroperoxyeicosatetraenoic acids (5S-, 12S-, 15S-HPETE), their reduced hydroxyl analogs, prostaglandins, and leukotriene B4 (Ahern 2003; Huang et al. 2006; Wang et al. 2005) (Fig. 2.1).



**Fig. 2.1** Selected examples of activators (1–8) and inhibitors (9–10) of TRPV1. **1** Olvanil (CID 5311093). **2** Piperine (CID 638024). **3** Allicin (CID 65036). **4** Anandamide (CID 5281969). **5** NADA: N-arachidonoyl dopamine. (CID 5282105). **6** OLEA: N-oleoyl ethanolamine (CID 5283454). **7** EMA-6: N-arachidonoyl serine (CID 10596625). **8** 15-HpETE: 15-hydroperoxy eicosatetraenoic acid (CID 6437084). **9** BCTC (CID 9929425). **10** I-RTX: 5-iodoresiniferatoxin (CID 16219535)

As expected, the activation of TRPV1 in nociceptors with vanilloids causes a burning pain sensation and irritation. Paradoxically, capsaicin has been in use for many years as anti-nociceptive compound in peripheral neuropathies (e.g., post-herpetic neuralgia, neuropathy, mastectomy, amputation and skin cancer). Capsaicin is used as an analgesic, because in addition to activate the channel, it also induces its desensitization. Furthermore, the repetitive application of the vanilloid produces a rundown of channel activity known as tachyphylaxia that results in a strong anti-nociceptive effect (Knotkova et al. 2008). This analgesia may be accompanied by reversible and/or irreversible loss of the capsaicin sensitive C-fibers (Hiura 2000).

Although TRPV1 agonists may have some therapeutic application, their low in vivo activity, along with their poor bioavailability and secondary effects has limited their development as anti-nociceptives, and promoted the research into the design of potent antagonists that display higher therapeutic index. The efforts in developing TRPV1 antagonists have been concentrated in obtaining both competitive and non-competitive (including uncompetitive) inhibitors (Planells-Cases et al. 2003; Szallasi and Appendino 2004). Uncompetitive antagonists acting as open channel

blockers are activity-dependent blockers that preferentially bind to over-activated receptors, with minimal interaction with the physiologically working channels. Accordingly, they are supposed to display lower side-effects than conventional antagonists.

Among the competitive TRPV1 antagonists (Fig. 2.1), capsazepine was the first identified, although with poor *in vivo* activity (Bevan et al. 1992; Walker et al. 2003). A vanilloid with better therapeutic potential is 5-iodo-RTX, a potent TRPV1 antagonist ( $IC_{50}=3.9$  nM) (McDonnell et al. 2002; Wahl et al. 2001). This compound produced notable analgesic activity *in vivo* and it is currently under clinical studies.

The family of competitive antagonists grew tremendously thanks to the contribution of pharmaceutical companies that established strong drug discovery programs for TRPV1 channels. As a result, ultra-high affinity synthetic antagonists were discovered for analgesic drug development. However, most of the clinical trials for these compounds had to be cancelled in Phase I because the indiscriminate blockade of TRPV1 channels with these compounds resulted in significant hyperthermia in humans, suggesting that this receptor also plays a pivotal role in core body temperature (Gavva et al. 2008).

The first non-competitive TRPV1 antagonist was the trinuclear polyamine complex, ruthenium red that was followed by arginine-rich peptides, and peptidomimetic compounds such as peptoids DD00069 and DD01050 (Garcia-Martinez et al. 2002, 2006). All these compounds resulted in unacceptable *in vivo* side effects and toxicity that prevented their clinical development. Recently, an uncompetitive antagonist, based in a triazine scaffold (triazine 8aA) that block TRPV1 channel by an activity-dependent mechanism was reported (Vidal-Mosquera et al. 2011). Triazine 8aA showed a strong voltage-dependent TRPV1 blockade by inhibiting at negative membrane potential, a hallmark of open-channel blockers. This compound holds promise for therapeutic development, although *in vivo* activity in pain models has not been yet reported.

Allosteric modulators of TRPV1 activity are another class of non-competitive antagonists. These compounds interfere with the allosteric mechanism that gates the channel. Structure-function analysis of TRPV1 channels demonstrated that the intracellular TRP domain, a highly conserved region adjacent to the receptor internal gate (Venkatachalam and Montell 2007), is essential for subunit tetramerization and allosteric activation (Garcia-Sanz et al. 2004, 2007). Thus, this protein interface could be used as an allosteric site to modulate channel function. Indeed, compound TRP-p5, a palmitoylated 13-mer peptide patterned after the N-terminus region of the TRP domain, displays *in vitro* and *in vivo* inhibitory activity (Valente et al. 2011). This finding is proof-of-concept that allosteric modulators such as TRPducins represent another family of non-competitive antagonists that could be developed therapeutically as anti-nociceptives.

A complementary approach to reduce the inflammatory sensitization of TRPV1 has been to interfere with the recruitment of the channel to the neuronal surface. This strategy has proven that blockers of neuronal exocytosis such as compound DD04107 display analgesic activity (Ponsati et al. 2012). *In vitro* experiments with DD04107 showed that it blocked the inflammatory over expression of TRPV1

channels to the plasma membrane (Camprubi-Robles et al. 2009). In vivo, this compound displays long-lasting anti-nociceptive activity against inflammatory and neuropathic pain, without apparent side effects, demonstrating that acting on the TRPV1 signalplex may be a valuable pharmacological strategy (Ponsati et al. 2012). This compound is being developed clinically.

## 2.3 TRPV2

At variance with TRPV1 channels, the pharmacology of its close homologue TRPV2 is still in its infancy (Peralvarez-Marin et al. 2013). This non-selective  $\text{Ca}^{2+}$  channel is also present in the peripheral nervous system and co-localizes with TRPV1 in a subset of nociceptors (Liapi and Wood 2005). The physiological role of this TRP channel is yet elusive. Initially was considered a thermoTRP channel that activated at  $52^\circ\text{C}$ , and also responded to hypotonicity (Caterina et al. 1999; Muraki et al. 2003). However, these are still highly debated functions (Park et al. 2011; Peralvarez-Marin et al. 2013), thus requiring further investigation, including the discovery of agonists and antagonists that could be used as pharmacological tools.

The identification of specific TRPV2 modulators is, surprisingly, inexistent, probably due to the species-specific pharmacology coupled with problems in developing stable recombinant cell lines due to cytotoxic effects of TRPV2 expression (Penna et al. 2006). Several chemical compounds have been shown to modulate TRPV2, however, virtually all of them are non-specific (Table 2.1 and Fig. 2.2). Indeed, TRPV2 is activated by general TRP channel agonists, such as 2-aminoe-thoxy-diphenyl borate (2-APB), probenecid, lysophospholipids, and cannabinoids (Juvin et al. 2007; Monet et al. 2009; Qin et al. 2008). However, the response to these ligands is low and variable and quite species-dependent (Neeper et al. 2007).

To date, only general blockers such as ruthenium red and trivalent cations ( $\text{La}^{3+}$  and  $\text{Gn}^{3+}$ ) (Table 2.1), have been described as blockers of TRPV2 (Lefler et al. 2007). In addition, the potassium channel blockers tetraethylammonium (TEA), 4-aminopyridine (4-AP), and 1-(2-(trifluoromethyl)phenyl)imidazole are also able to block TRPV2 currents (Vriens et al. 2009). Other reported inhibitors are SKF96365, amiloride, and Tranilast, an antiallergic drug (Juvin et al. 2007; Mihara et al. 2010) (Fig. 2.2).

## 2.4 TRPV3

TRPV3 is a non-selective  $\text{Ca}^{2+}$  channel that plays a pivotal role in various physiological processes in the skin and hair follicles. This channel displays a moderate sequence homology to TRPV1. TRPV3 is mainly located in keratinocytes and epithelial cells (Nilius and Owsianik 2011; Valdes-Rodriguez et al. 2013), and marginally in sensory neurons (Nilius et al. 2014). This TRP channel is a polymodal receptor

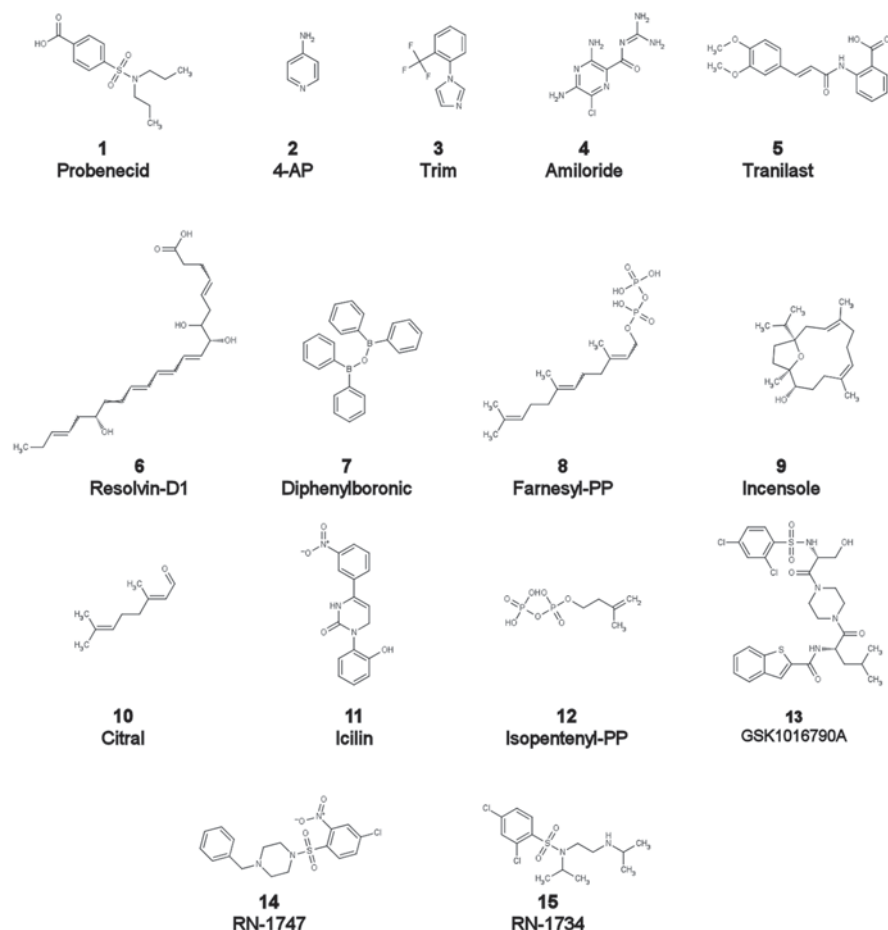


**Table 2.1** Representative modulators of depicted TRP channels

Ion channel	Activators	Representative blockers
TRPV1	Capsaicin, resiniferatoxin, olvanil, piperine, eugenol, camphor, 2-APB, allicin, anandamide, NADA, OLEA, N-arachidonolylserine 5S-, 12S-, 15S-HPETE, prostagalandine, leukotriene B4	Capsazepine, ruthenium red, DD01050, 5-iodo-RTX, Triazine 8aA, TRP-p5
TRPV2	2-APB, probenecid, lysophospholipids, cannabinoids	Ruthenium red, La <sup>3+</sup> , Gn <sup>3+</sup> , TEA, 4-aminopyridine, 1-(2-(trifluoromethyl)phenyl) imidazole, SKF96365, amiloride, Tranilast.
TRPV3	2-APB, 17(R)-resolvin D1, PIP <sub>2</sub> , diphenylboronic anhydride, farnesyl pyrophosphate camphor, carvacrol, eugenol, menthol, thymol, borneol, cresol, carveol, gerianool, propofol, linalool, incensole, citral	Ruthenium red, icilin, isopentenyl pyrophosphate chromane-, fused pyrimidine-, fused pyrimidinones-, chromanone- and fused imidazole-derivatives
TRPV4	Endocannabinoids, arachidonic acid metabolites, nitric oxide, diacylglycerol, bisandrographolide A, 4 $\alpha$ PDD phorbol derivatives, GSK1016790A, RN-1747	RN-1734
TRPC5	Thioredoxin, lysophosphatidylcholine, lanthanides, genistein, diadzein	SKF-96365, BTP-2, flufenamic acid, chlorpromazine, W-13, calmidazolium, W-7, 2-APB, ML-7, ML-9
TRPM3	Pregnenolone sulphate, dihydro-D-erythro-sphingosine, N,N-dimethyl-D-erythro-sphingosine, dihydropyridine nifedipine	2-APB, Gd <sup>3+</sup> , rosiglitazone, troglitazone, mefenamic acid, cholesterol, naringenin, hesperetin
TRPM8	Menthol, icilin, geraniol, D3263	AMTB, JNJ41876666, BCTC, Thio-BCTC, clotrimazole, econazole, SKF-96365
TRPA1	Allyl isocyanate, cinnamaldehyde, allicin, nifedipine, chlorpromazine, auranofin, clotrimazole, clioquinol, apomorphine, glibenclamide, BCTC	HC-030031, GRC-17536, A-967079, piperazineurea, N-1-Alkyl-2-oxo-2-aryl amide, 1,8-cineole, chlorpromazine, toxin ProTx-I

activated by non-painful temperatures (Peier et al. 2002a; Smith et al. 2002; Xu et al. 2002), and chemical stimuli (Xu et al. 2006a), including natural irritants and synthetic ligands (Xu et al. 2006a), and endogenous compounds, some of them involved in the downstream inflammatory cascade (Doerner et al. 2011; Sherkheli et al. 2009). Stimulation of TRPV3 releases inflammatory mediators from keratinocytes including ATP, prostaglandin E2 and IL-1, which supports its contribution to pain transduction and inflammatory signaling. Indeed, in certain human disease states there are changes in the expression of TRPV3, such as an increase in painful breast tissue (Matta et al. 2008), or a decrease in keratinocytes in diabetic neuropathy (Facer et al. 2007).

Some evidence points to phosphatidyl inositol-4,5-bisphosphate and 17(R)-resolvin D1 as putative *in vivo* modulators of TRPV3 (Bang et al. 2012; Doerner et al. 2011) (Table 2.1). A role of 17(R)-resolvin D1 as potential analgesic mediated by TRPV3 has been described, although a direct evidence is still missing (Bang et al.



**Fig. 2.2** Selected examples of TRPV2-4 effectors. TRPV2 activators (1) and inhibitors (2–5). TRPV3 activators (6–10) and inhibitors (11–12). TRPV4 activators (13–14) and inhibitors (15). **1** probenecid (CID 4911). **2** 4-AP: 4-aminopyridine (CID 1727). **3** Trim: 1-(2-(trifluoromethyl)phenyl)imidazole (CID 1359). **4** Amiloride (CID 16231). **5** Tranilast (CID 5282230). **6** Resolvin-D1: 17(R)-resolvin D1 (CID 71434077). **7** Diphenylboronic anhydride (CID 596810). **8** Farnesyl pyrophosphate (CID 44134714). **9** Incensole (CID 44583885). **10** Citral (CID 638011). **11** Icilin (CID 161930). **12** Isopentenyl pyrophosphate (CID 1195). **13** GSK1016790A (CID 23630424). **14** RN-1747 (CID 5068295). **15** RN-1734 (CID 3601086)

2012). Similarly, TRPV3 has been related with the production of nitric oxide via a nitrite independent pathway (Miyamoto et al. 2011). Furthermore, farnesyl pyrophosphate and isopentenyl pyrophosphate, intermediates of the melanovate pathway, are activator and inhibitor respectively, suggesting a fine-tuning of TRPV3

function (Bang et al. 2010; 2011). Alfa-hydroxy acids are proton donors commonly used in cosmetics to produce skin exfoliation mediated by TRPV3 activation (Cao et al. 2012).

The pharmacology of the TRPV family is far from simple, and TRPV3 is not an exception (Table 2.1 and Fig. 2.2). 2-APB also activates TRPV3 (Chung et al. 2004; Hu et al. 2004, 2009). Prolonged exposure of TRPV3 to 2-APB induced sensitization (Sherkheli et al. 2009). Structurally related 2-APB compounds such as diphenylboronic anhydride also act as potent TRPV3 agonists (Chung et al. 2005).

Natural aromatic monoterpenes, such as camphor, carvacrol, eugenol, menthol, thymol, as well as borneol, cresol, and others are an additional class of TRPV3 ligands (Moqrich et al. 2005; Vriens et al. 2009; Xu et al. 2006a). Camphor is a weak agonist for TRPV3 that activates currents only at concentrations of 10 mM. Carvacrol is responsible for arterial vasodilation by activating TRPV3 channels in the endothelium (Earley et al. 2010), which may account for some of their attributed cardioprotective effects. In addition to camphor and carvacrol, thymol and eugenol have also been shown to enhance the temperature response of TRPV3 (Macpherson et al. 2006; Xu et al. 2006a).

Non-aromatic monoterpenes such as carveol and derivatives (monocyclic), or geraniol, propofol and linalool (acyclic) display strong TRPV3 agonism (Vogt-Eisele et al. 2007). Incensole and incensole acetate are diterpenic cembrenoids found in incense (*Boswellia papyrifera*) potently activate TRPV3. The traditional use of these natural products is related to anti-inflammatory effects through the activation of TRPV3 in the skin. Interestingly, incensole acetate produces anxiolytic and antidepressive effects in mice (Moussaieff and Mechoulam 2009; Paul and Jauch 2012). Citral, a bioactive component of lemongrass is also an agonist of TRPV3 (Stotz et al. 2008), adding to the list of compounds acting on this channel (Fig. 2.2).

Cannabinoids such as cannabidiol or delta-9-tetrahydrocannabinol modulate nonspecifically TRPV3. Other derivatives such as cannabigerovarin or cannabigerolic desensitize TRPV3 (De Petrocellis et al. 2012). Active research is necessary in this field because, interestingly, the activation of TRPV3 by these compounds may contribute to their described *in vivo* activity (Anand 2003; Galeotti et al. 2001; Santos and Rao 2001; Umezu et al. 2001; Xu et al. 2005a).

TRPV3 antagonists include the non-specific ruthenium red, that blocks all TRPV family member at negative potentials (Vennekens et al. 2008). The compound icilin, which is a strong agonist of TRPM8 channel, is an inhibitor of TRPV3 at low doses (Sherkheli et al. 2012). Novel inhibitors are under study and have promising analgesic effects, which further suggests the involvement of TRPV3 in pain transduction (Reilly and Kym 2011). Several pharmaceutical industries have reported strong and selective TRPV3 antagonists including series of chromane-, fused pyrimidine-, fused pyrimidinones-, chromanone- and fused imidazole-derivatives (Ferrer-Montiel et al. 2012). Some of these antagonists are currently under clinical studies to treat human pain conditions.

## 2.5 TRPV4

Transient Receptor Potential Vanilloid 4 (TRPV4) a non-selective  $\text{Ca}^{2+}$  channel is a homologue of the OSM-9 osmosensory channel first described in *C. elegans*. TRPV4 is activated by warm temperatures (27–35 °C) (Guler et al. 2002; Liedtke et al. 2000), and is sensitive to cell swelling and shear stress (Gao et al. 2003; Kohler et al. 2006; Loukin et al. 2010; Strotmann et al. 2000). Functions include temperature monitoring in skin keratinocytes, osmolarity sensing in the kidney (Pochynyuk et al. 2013), and shear stress detection in blood vessels, which indicates that TRPV4 functions as a putative mechanosensor (Nilius et al. 2003a, b), and is involved in nociception (Alessandri-Haber et al. 2005, 2006). It has been reported that TRPV4 may be activated by hypotonic solutions, and by mechanical forces in membrane patches (Loukin et al. 2009). TRPV4 may contribute to development of mechanical hyperalgesia after inflammation and injury (Alessandri-Haber et al. 2006). This channel is expressed in several tissues, including primary sensory neurons (Alvarez et al. 2006; Birder et al. 2007; Guler et al. 2002; Pochynyuk et al. 2013; Strotmann et al. 2000; Tabuchi et al. 2005; Watanabe et al. 2002b; Yang et al. 2006).

TRPV4 is activated by endogenous chemical ligands, such as endocannabinoids, arachidonic acid metabolites and nitric oxide (Birder et al. 2007) (Table 2.1 and Fig. 2.2). Phorbol esters that do not activate PKC, mediate TRPV4 heat responses (Watanabe et al. 2002a). TRPV4 sensitivity to osmotic and mechanical stimuli may depend on phospholipase A2 activation and the generation of arachidonic acid metabolites (Fernandes et al. 2008; Liedtke et al. 2000; Strotmann et al. 2000; Vriens et al. 2004). Furthermore, TRPV4 is activated by hypotonicity, diacylglycerol, and PKC-activating phorbol esters (Watanabe et al. 2002a, b, 2003).

Natural plant extracts (Klausen et al. 2009), bisandrographolide A (Smith et al. 2006) and synthetic compounds, such as a phorbol derivative (Birder et al. 2007), or GSK1016790A (Thorneloe et al. 2008) also activate TRPV4 channels. In addition, small molecules such as compound RN-1747 was also found to be a TRPV4 agonist (Vincent et al. 2009) (Table 2.1 and Fig. 2.2).

TRPV4 antagonism is being considered for inflammatory and neuropathic pain treatment (Vincent and Dunton 2011). However, selective TRPV4 antagonists have not been described appropriately. Ventilator-induced lung injury has emerged as a potential indicator for TRPV4 antagonists (Jin et al. 2011) (Table 2.1). The small molecule RN-1734 2,4-Dichloro-N-isopropyl-N-(2-isopropylaminoethyl)benzenesulfonamide was observed to inhibit ligand- and hypotonicity-activated TRPV4 (Vincent et al. 2009). In addition, the compound showed selective properties for TRPV4 over other TRPs such as TRPV1, TRPV3 and TRPM8, being a valuable pharmacological tool for TRPV4 studies (Vincent et al. 2009).

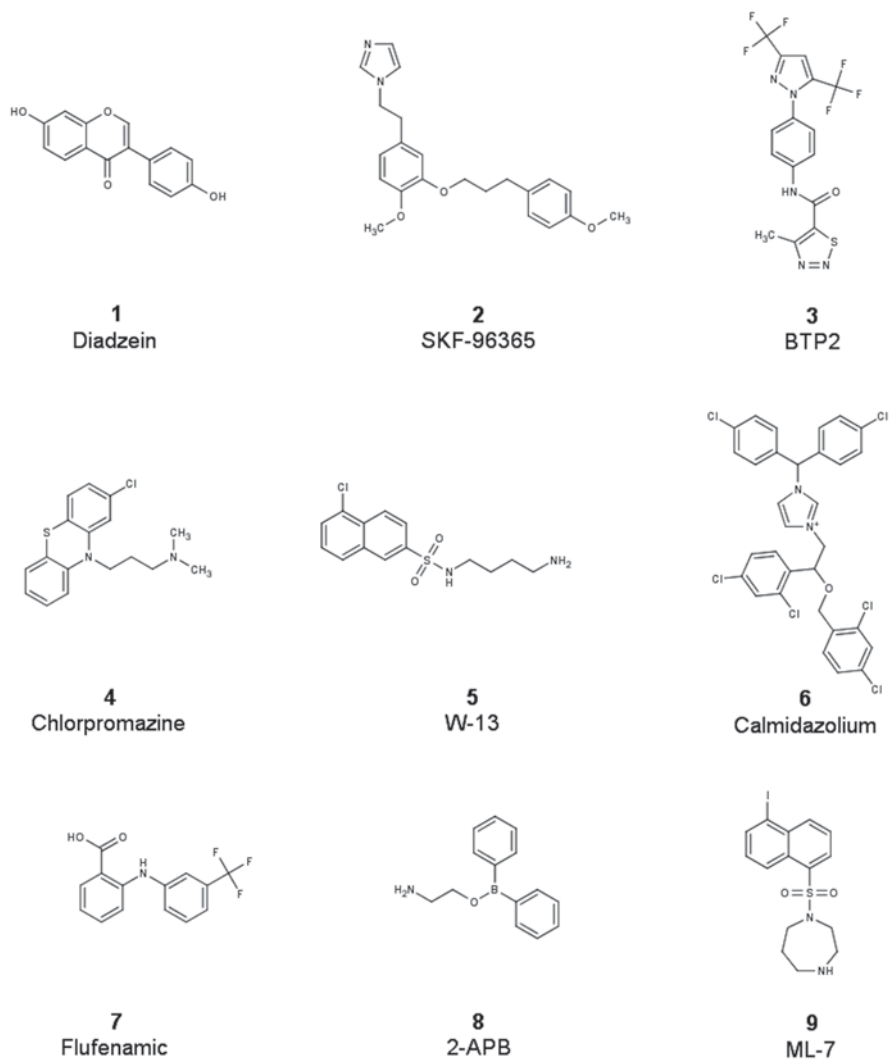
## 2.6 TRPC5

Several mammalian and *Drosophila* TRP canonical, TRPC proteins (TRPC1-7) have been identified (Plant and Schaefer 2003; Wes et al. 1995; Hardie and Minke 1995). All mammalian TRPCs seem to be enhanced with G-protein-coupled receptors and tyrosine kinases receptors (Montell 1999). The channels may be divided in three subgroups according to sequence homology: C1-C4-C5, C3-C6-C7, and C2 (Zufall et al. 2005). Particularly, TRPC5 is a functional plasma membrane ion channel (Beech 2007) activated by hypo-osmotic stimuli, which is dependent on phosphoinositides (Gomis et al. 2008). The inhibition of TRPC5 has been shown to suppress inflammatory pain induced by the component of the bee venom mellitin (Ding et al. 2011). Several studies support the conclusion that TRPC5 plays a role in growth cone extension and axonal guidance (Davare et al. 2009). A variety of other functions have been assigned to TRPC5, indicating a central role of this channel in physiology (Jiang et al. 2011; Nath et al. 2009; Premkumar and Abooj 2013; Wu et al. 2010; Wuensch et al. 2010; Xu et al. 2008), although the channel is not essential for life (Ricchio et al. 2009).

TRPC5 modulation is not well known. A common stimulus for TRPC5 is the activation of G protein-coupled receptor. Many different receptors may be involved, including receptors for adenosine 5'-triphosphate, bradykinin, acetylcholine, histamine, prostaglandin E2, thrombin, uridine 5'-triphosphate, sphingosine-1-phosphate, glutamate and cholecystokinin (Meis et al. 2007; Ricchio et al. 2009; Xu et al. 2006b; Zeng et al. 2004). TRPC5 is stimulated by activation of growth factor receptors (Bezzarides et al. 2004). TRPC5 is also a target for thioredoxin, an endogenous redox protein with established intracellular functions. Reduced thioredoxin activates TRPC5 expressed in secretory fibroblast-like synoviocytes when secreted extracellularly in patients with rheumatoid arthritis (Xu et al. 2008). Lysophosphatidylcholine has been identified as a TRPC5 activator (Flemming et al. 2006). Current data suggest a complex arrangement between TRPC5 activity and various lipid factors, supporting the hypothesis that a physiological function of TRPC5 channels is to act as lipid signal transducers.

An unusual feature of TRPC5 is its stimulation by external lanthanides (Jung et al. 2003; Schaefer et al. 2000; Xu et al. 2005b; Zeng et al. 2004). It was reported that ionic lead ( $Pb^{2+}$ ) mimics the effect of lanthanides, leading to the hypothesis that TRPC5 may confer survival advantage by acting as a sensor of heavy metal ions (Sukumar and Beech 2010). Stimulation of TRPC5 by isoflavones like genistein or diadzein has also been reported (Wong et al. 2010) (Table 2.1, Fig. 2.3).

Although no specific or potent exogenous chemical inhibitors of TRPC5 are known, various chemicals have effects on TRPC5 function (Table 2.1). In many of these cases, it is not clear if the agent acts directly on the channel. TRPC5 has been reported to be inhibited by SKF-96365 (Okada et al. 1998), 3,5-bis(tri-fluoromethyl)pyrazole derivative BTP-2 (He et al. 2005; Kiyonaka et al. 2009), flufenamic acid (Lee et al. 2003b), W-13 or chlorpromazine (Shimizu et al. 2006), W-7 or calmidazolium (Kim et al. 2006), Pyr2, 2-APB (Xu et al. 2005b), and the myosin light chain kinase inhibitors ML-7 or ML-9 (Shimizu et al. 2006) (Fig. 2.3).



**Fig. 2.3** Selected examples of activators (1) and inhibitors (2–9) of TRPC5. **1** Diadzein (CID 5281708). **2** SKF-96365 (CID 11957693). **3** BTP2 (CID 2455). **4** Flufenamic acid (CID 3371). **5** Chlorpromazine (CID 2726). **6** W-13 (CID 4299). **7** Calmidazolium (CID 644274). **8** 2-APB (CID 1598). **9** ML-7 (CID 4216)

## 2.7 TRPM3

TRPM3, TRP melastatin 3, non-selective  $\text{Ca}^{2+}$  channel is one of the least investigated proteins of the TRP family of ion channels. In humans, it is highly expressed in the kidney (Grimm et al. 2003; Lee et al. 2003a), brain (Lee et al. 2003a; Oberwinkler 2007), sensory neurons, human pituitary (Fonfria et al. 2006), vascular smooth

muscle (Naylor et al. 2010) and pancreatic beta cells (Thiel et al. 2013; Wagner et al. 2008). However, its physiological role is still under investigation. Activation of TRPM3 has been linked to insulin secretion in pancreatic beta-cells (Wagner et al. 2008), to vascular smooth muscle cell contraction (Naylor et al. 2010), and to potentiating glutamatergic transmission in cerebellar Purkinje neurons of developing rats (Zamudio-Bulcock et al. 2011). A role in pain transduction has also been reported for TRPM3 (Vriens et al. 2011).

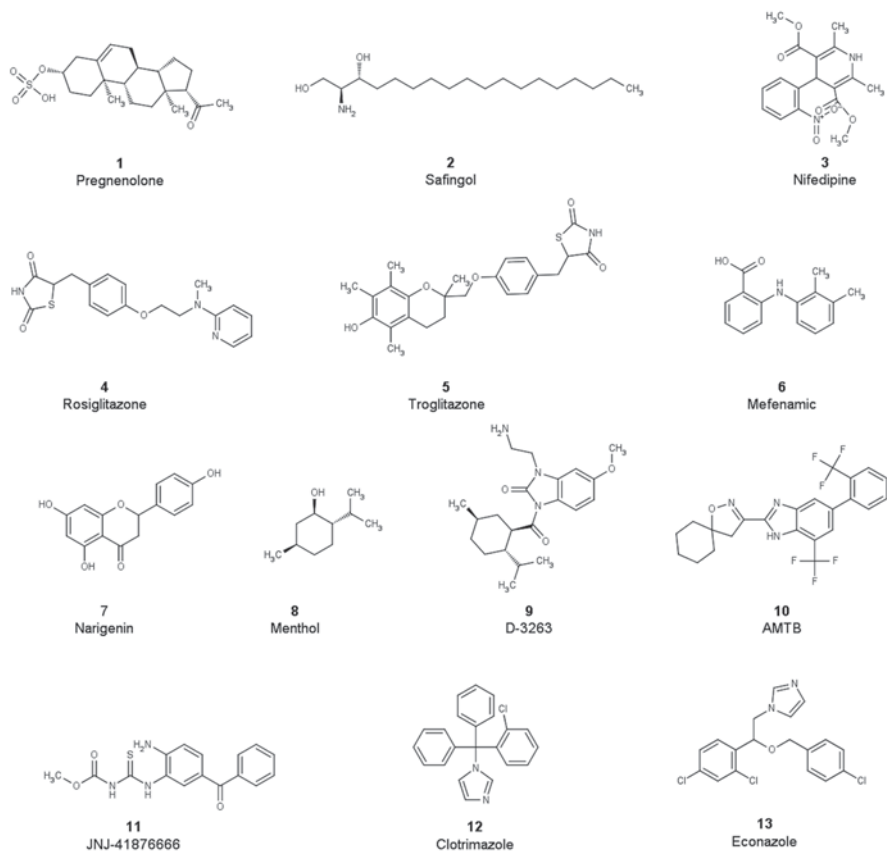
Recently, pharmacological investigations have been initiated in order to identify substances that influence TRPM3 channel activity. TRPM3 is rapidly and reversibly activated by extracellular pregnenolone sulphate, a neuroactive steroid. Application of pregnenolone sulphate led to a rapid calcium influx and enhanced insulin secretion from pancreatic islets (Wagner et al. 2008). Pregnenolone sulfate also activates TRPM3 channels in HEK293 cells, vascular smooth muscle cells, and synovial fibroblasts (Ciurtin et al. 2010; Klose et al. 2011; Majeed et al. 2012; Naylor et al. 2010), confirming the functional relevance of TRPM3 in contractile function. However, the concentration of pregnenolone sulfate required to stimulate TRPM3 channels is in the micromolar range, suggesting that pregnenolone sulfate is not a physiological agonist of TRPM3 and may have only pharmacological relevance. Moreover, the fact that TRPM3 deficient mice did not show alterations in resting blood glucose levels (Vriens et al. 2011) suggests that TRPM3 plays a marginal role in controlling  $\beta$ -cell functions.

Two structural analogs of sphingosine, dihydro-D-erythro-sphingosine and N, N-dimethyl-D-erythro-sphingosine, are able to activate TRPM3 (Grimm et al. 2005) (Table 2.1 and Fig. 2.4). Surprisingly, TRPM3 channels are also activated by the dihydropyridine nifedipine, an inhibitor of voltage-gated  $\text{Ca}^{2+}$  channels, while the structurally related compounds nimodipine, nicardipine, and nitrendipine were inactive (Wagner et al. 2008).

As for many other members of the TRP ion channel family, 2-APB and  $\text{Gd}^{3+}$  have been reported to inhibit  $\text{Ca}^{2+}$  influx through TRPM3 channels (Grimm et al. 2003; Harteneck and Schultz 2007; Xu et al. 2005b). Other TRPM3 channel blockers described thus far include the antidiabetic PPAR $\gamma$ -agonists rosiglitazone and troglitazone (Majeed et al. 2012). Nonsteroidal anti-inflammatory drugs (NSAIDs) of the fenamate group, like mefenamic acid are able to selectively block TRPM3-mediated  $\text{Ca}^{2+}$  entry as well as insulin release (Klose et al. 2011). Cholesterol, the precursor metabolite of pregnenolone and progesterone, also prevents TRPM3 channel activation. (Naylor et al. 2010).

Recently, the screening of a compound library revealed that citrus fruit flavanones, such as naringenin and hesperetin, and fabacea secondary metabolites selectively inhibit TRPM3 channel activation with potencies ranged from upper nanomolar to lower micromolar concentrations (Straub et al. 2013) (Table 2.1).





**Fig. 2.4** Selected examples of TRPM3 and TRPM8 effectors. TRPM3 activators (1–3) and inhibitors (4–7). TRPM8 activators (8–9) and inhibitors (10–13). **1** Pregnenolone sulphate (CID 105074). **2** Safingol: Dihydro-D-erythro-sphingosine (CID 91486). **3** Nifedipine (CID 4485). **4** Rosiglitazone (CID 77999). **5** Troglitazone (CID 5591). **6** Mefenamic acid (CID 4044). **7** Narigenin (CID 932). **8** Menthol (CID 16666). **9** D-3263 (CID 44137358). **10** AMTB (CID 3036972). **11** JNJ-41876666: Johnson&Johnson patent. **12** Clotrimazole (CID 2812). **13** Econazole (CID 3198)

## 2.8 TRPM8

TRPM8 is a  $\text{Ca}^{2+}$  permeable channel that was first identified in prostate cancer cells (Tsavalier et al. 2001; Bidaux et al. 2007), but also is present along the male urogenital tract (De Blas et al. 2009), artery myocytes (Johnson et al. 2009), and lung epithemium cells (Sabnis et al. 2008), although its role in many of these tissues still remains unclear. These channels are expressed in primary sensory neurons in skin and mucosae, with a physiological role in detecting low temperature signals (10–33 °C) (Babes et al. 2011), and in sensing cooling chemicals like menthol and icilin (McKemy et al. 2002; Peier et al. 2002) (Bharate and Bharate 2012; Chuang

et al. 2004). Under pathological conditions, there are increasing experimental evidences that confirm the anomalous over-expression of TRPM8 channels in sensory neurons after nerve injury or inflammation, as well as their involvement in cold allodynia and hyperalgesia (Kapoor 2012; Xing et al. 2007) (Abe et al. 2006; Ramachandran et al. 2013). The activation of TRPM8 also attenuates pain in certain acute and inflammatory pain states, mediating for instance the analgesic effects of menthol (Liu et al. 2013). Therefore, both TRPM8 agonists and antagonists could be valuable analgesic agents (Liu and Qin 2011; Maelkiaie et al. 2011). TRPM8 channels are also expressed in corneal afferent neurons implicated in the regulation of ocular surface wetness, and in this respect TRPM8 modulators could have application in dry eye syndrome and excessive lacrimation dysfunction (Fernández-Peña and Viana 2013; Parra et al. 2010).

On the other hand, TRPM8 is abnormally over-expressed in androgen-sensitive prostate cancer (Gkika and Prevarskaya 2011; Tsvaler et al. 2001), breast cancer (Dhennin-Duthille et al. 2011; Ouadid-Ahidouch et al. 2012), skin melanoma cells (Yamamura et al. 2008), human pancreatic adenocarcinoma (Yee et al. 2010), oral scamous cell carcinoma (Okamoto et al. 2012), and osteosarcoma tissues and cell lines (Wang et al. 2014). Again, both agonists and antagonists of TRPM8 have proved to be valid as pharmacological tools for reducing growth and progression of neoplasias with intense expression of TRPM8 channels. Therefore, TRPM8 may also be considered an attractive target for therapeutic intervention in the search for new antitumor agents (Knowlton and McKemy 2011; Lehen'kyi and Prevarskaya 2011).

The crucial role of TRPM8 in the human pathologies is behind the intensive drug-discovery programs developed in recent years around this channel (with more than 25 patents filed since 2009). Among the antagonists, different families having benzothiophene, benzimidazole and arylglycine moieties as the central scaffold have been reported (Calvo et al. 2012; Matthews et al. 2012; Parks et al. 2011; Zhu et al. 2013). Some of these compounds showed excellent *in vitro* and *in vivo* profiles, including activity in animal models of inflammatory and neuropathic pain, thus emerging as strong candidates for future development (Table 2.1). The benzothiophene derivative JNJ41876666 has been used, along with other antagonist, AMTB, and RNAi to determine that the inhibition of either the expression or function of TRPM8 channels reduces the proliferation rate of prostate tumor cells, while no effect was observed in non-tumor cells (Valero et al. 2012).

The commercial antagonist N-(3-Aminopropyl)-2-[(3-methylphenyl)methoxy]-N-(2-thienylmethyl) benzamide hydrochloride AMTB has also served to suggest the potential of TRPM8 channel blocker as a new therapeutic opportunity for treating overactive bladder and painful bladder syndrome (Lashinger et al. 2008). Some tetrahydroquinoline and aza-analogues are TRPM8 antagonists, and a selected compound from this series reduced icilin-induced wet-dog shakes (WDS) in a dose dependent manner (Tamayo et al. 2012). Other chemotypes able to inhibit TRPM8 channels include the piperazine urea derivative BCTC, used to demonstrate that the menthol- and cold-induced allergic responses of mast cells are mediated by TRPM8 (Cho et al. 2010). A series of spiro-chromene-piperidines endowed with high po-

tency and favorable ADME properties are effective in rodent models of neuropathic pain (Chaudhari et al. 2013). SKF-96365 and the antifungal drugs clotrimazole and econazole are also effective blockers of TRPM8 (Madrid et al. 2006; Mälkiä et al. 2009 Meseguer et al. 2014; Table 2.1 and Fig. 2.4).

It has also been described that the Gαq protein, formed after activation of GPCRs, blocks TRPM8 activity by the direct formation of a complex with the channel (Zhang et al. 2012). This could indirectly arbitrate the inhibition of TRPM8 by the inflammatory mediators bradykinin and histamine in sensory nerves, and could open new strategies for modulating these channels, interfering within protein-protein interactions.

TRPM8 agonists are also therapeutically important for attenuating pain, and may induce apoptosis in TRPM8 expressing cancer cells (Maelkiae et al. 2011) (Table 2.1). The most remarkable result in this field refers to the ability of 3-(2-aminoethyl)-1(R)-[(2(S)-isopropyl-5(R)-methyl cyclohexanecarbonyl)]-5-methoxy-1,3-dihydro-benzimidazol-2-1 hydrochloride D3263 to inhibit the growth of TRPM8 expressing tumors. This new orally bioavailable chemical entity has already completed Phase I clinical trials in healthy individuals. The company also reported clinical studies on patients with advanced solid tumors, for which the preliminary results indicate disease stabilization after treatment. Some preclinical data also demonstrated the potential of D3263 to treat benign prostatic hyperplasia by itself or in combination with the synthetic 5α-reductase inhibitor finasteride.

## 2.9 TRPA1

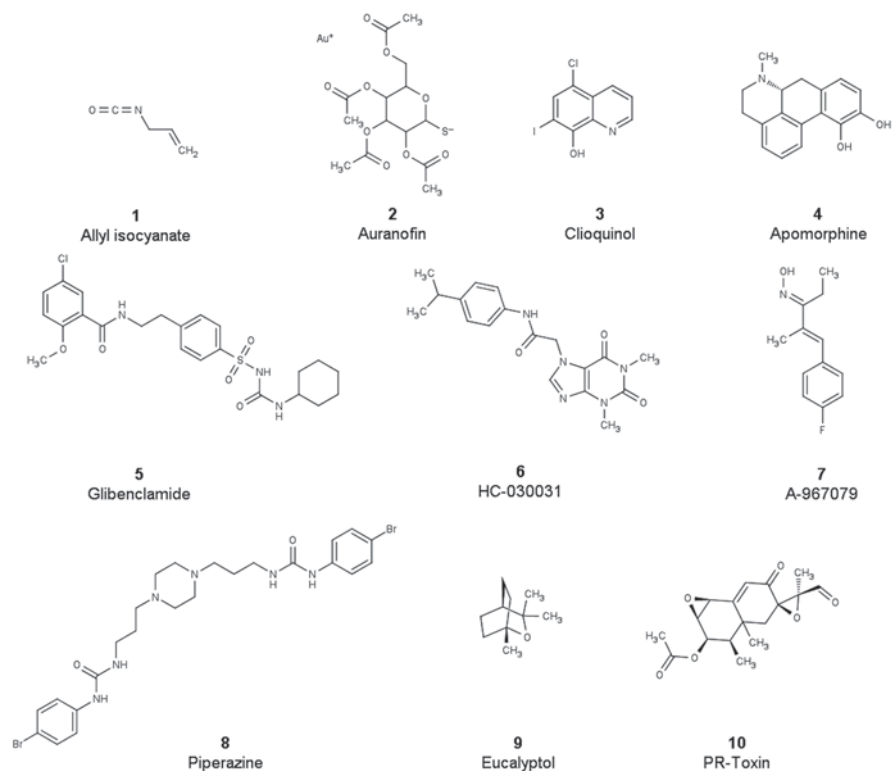
Transient receptor potential ankyrin 1 (TRPA1) is a non-selective Ca<sup>2+</sup> channel characterized by a high number of ankyrin repeats (14) at the N-terminal domain (Andrade et al. 2012). This channel is activated by multiple stimuli, including temperature, acids, and numerous chemicals, like different noxious environmental and industrial pollutants, oxidant agents and bacterial endotoxins (Bandell et al. 2004; Bautista et al. 2005; Jordt et al. 2004; Levine and Alessandri-Haber 2007; Peterlin et al. 2007; Meseguer et al. 2014). Experimental evidence suggests that TRPA1 is activated by low temperatures, near the threshold of harmful cold for humans (Abrahamsen et al. 2008; del Camino et al. 2010; Karashima et al. 2009), although the categorization of TRPA1 as a cold thermosensor has been controversial.

TRPA1 receptors are vastly expressed in different unmyelinated sensory neurons (Story et al. 2003). They are also expressed in different organs, including the cardiovascular, gastrointestinal, and urinary systems (Andrade et al. 2012). TRPA1 co-localizes with TRPV1 channels at least in a subpopulation of C-type nociceptors (Fajardo et al. 2008). Akin to TRPV1, a number of studies have also established a crucial role of TRPA1 channels in neuronal and non-neuronal neuropathic pain (Barriere et al. 2012; Chen et al. 2011; Wei et al. 2010).

TRPA1 is also involved in respiratory reflexes due to inhaled pollutants (Bessac et al. 2009). These results suggest that the activation of this channel might con-

tribute to asthma and chronic obstructive pulmonary disease (Andre et al. 2008; Materazzi et al. 2010). TRPA1 is also activated by electrophilic products generated during oxidative processes, suggesting that these channels may act as sensors for the tissue damage during inflammatory processes (Taylor-Clark et al. 2008). Furthermore, TRPA1 channels have been identified as targets of allyl isocyanate, cinnamaldehyde, nifedipine, chlorpromazine, auranofin, as well as of clotrimazole, clioquinol, apomorphine, glibenclamide (Andrade et al. 2012; Babes et al. 2013), which provides some pharmacological basis for painful side effects of some of these drugs (Table 2.1 and Fig. 2.5).

The TRPA1 channel is considered a promising target for the development of new, clinically relevant drugs in different therapeutic areas (Baraldi et al. 2010). In this respect, the past few years have seen the emergence of novel TRPA1 antagonists, with more than 30 patents filed by different academic and non-academic institutions. Compounds claimed in these patents belong to different chemical



**Fig. 2.5** Selected examples of activators (1–5) and inhibitors (6–10) of TRPA1. 1 Allyl isocyanate (CID15123). 2 Auranofin triethylphosphane (CID 24199313). 3 Clioquinol (CID 2788). 4 Apomorphine (CID 6005). 5 Glibenclamide (CID 3488). 6 HC-030031 (CID 1150897). 7 A-967079 (CID 42641861). 8 Piperazine urea (CID 96934). 9 Eucalyptol: 1,8-cineole (CID 2758). 10 PR-Toxin (CID 56844124)

families, but a number of them are related to fused pyrimidindione derivatives displaying efficacy in various models of pain (Table 2.1). Among the pyrimidindione antagonists, we can mention compound HC-030031 and its isobutyl analogue Chembridge-5861528, which are being profusely used as pharmacological tools for studying the implication of TRPA1 channels in pathophysiological pain, among other disorders (Koivisto et al. 2012; Meotti et al. 2013; Samer et al. 2008; Shigetomi et al. 2012). A small library of related pyrrolo[3,2-d]pyrimidinone derivatives with micromolar antagonist potencies was described (Baraldi et al. 2012). In addition, compound GRC-17536 has successfully completed Phase I clinical trials and, since 2012, is under Phase II studies in patients with painful diabetic neuropathy. Good efficacy was also observed when this selective compound was administered by the inhalation route, and a Phase IIa study is ongoing in people with refractory chronic cough.

The oxime derivative A-967079 is a TRPA1 antagonist that inhibits  $\text{Ca}^{2+}$  influx through this channel at nanomolar concentrations, displayed moderate/good oral bioavailability, and was active in models of inflammatory and neuropathic pain (Chen et al. 2011). Related analogues, showing either modest TRPA1 agonist or antagonist properties, have been described (DeFalco et al. 2010). Other chemotypes displaying potent TRPA1 antagonist properties include N-arylsulfonyl-proline derivatives, piperazineurea, and N-1-Alkyl-2-oxo-2-aryl amide (Vallin et al. 2012). In addition, a series of 3-ylidenephthalides and some leucettamol marine products were described to display a dual action, since they are able to activate the TRPA1 channel and to block the related cold sensor TRPM8 (Chianese et al. 2012; Ortar et al. 2013). On the contrary, the 1,8-cineole, an essential oil from eucalyptus, evoked inward currents through human TRPM8, but inhibited TRPA1 activation (Takaishi et al. 2012). BCTC, a good blocker of TRPM8 at micromolar concentrations is also a strong activator of TRPA1 at similar concentrations (Madrid et al. 2006) (Table 2.1 and Fig. 2.5). All these structures may be considered versatile templates towards novel TRP channel modulators.

Very recently, toxin ProTx-I was identified as a high-affinity TRPA1 antagonist. This Cys-rich peptide, isolated from the venom of the Peruvian green-velvet tarantula, also behave as an antagonist of voltage-gated sodium ( $\text{NaV}1.2$ ) channels (Gui et al. 2014). However, mutations by Ala-scan indicated subtle differences in the structural requirements for binding to both ion channels. These findings open the possibility of using this peptide as the starting point for the development of new TRPA1 modulators. The use of ProTx-I, its mutants and the above indicated antagonists as pharmacological tools could certainly contribute to deepen our knowledge of TRPA1 function under physiological and pathological conditions, and to shed light on its gating mechanism.

## 2.10 Concluding Remarks

TRP channels pharmacology has been evolving a different pace for the members of this channel family, resulting in a plethora of modulators for few TRP proteins and a lack of ligands for the vast majority. However, technical advances in automated electrophysiology, along with an increase in the chemical diversity provided by synthetic and natural libraries most likely will change this conspicuous pharmacological unbalance. Furthermore, the discovery that allosteric modulators may be derived from the protein sequence will probably accelerate the discovery of TRP modulators. Moreover, the finding that TRP channel signalplexes are central to their function open new venues for drug intervention by targeting protein complexes involved in channel expression or signaling. Taken together, all these strategies will expand TRP pharmacology, even to previously considered non-druggable TRP channels.

**Acknowledgement** We are indebted to the members of our laboratories, to our collaborators, and the funding from the Ministry of Economy and Competitiveness (BFU2012-39092-C02-01/02, and CSD2008-00005), and the Generalitat Valenciana (PROMETEO/2010/046 and ISIC/2012/009).

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# Chapter 3

## Modulation of TRP Channels by N-glycosylation and Phosphorylation

**María Pertusa and Rodolfo Madrid**

**Abstract** Transient receptor potential (TRP) are cation permeable channels found in eukaryotic organisms, from yeast to human. These channels are involved in several physiological processes ranging from  $\text{Ca}^{2+}$  and  $\text{Mg}^{2+}$  homeostasis to sensory functions, including vision, pheromone signalling, taste, nociception, mechanotransduction and temperature sensing, among others. Most of these channels show covalent post-translational modifications (PTMs) that have a profound effect on their physiological function, by influencing their subcellular trafficking and/or their biophysical properties. Among these modifications, N-linked glycosylation and phosphorylation are two of the most relevant PTMs determining the functional expression of TRP channels. In this chapter we summarize their role in modulating the functional properties of TRP channels, with a critical role in sensory transduction.

**Keywords** N-glycosylation · Phosphorylation · Post-translational modifications · TRP channels

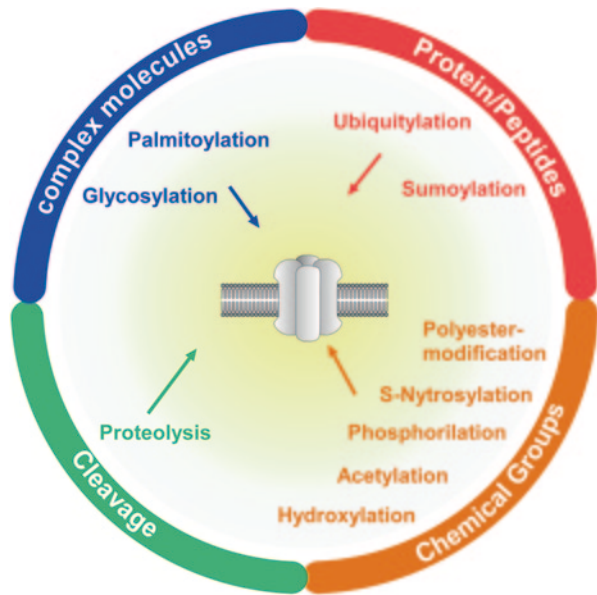
### 3.1 Introduction

In eukaryotic cells, proteins can be modified after translation by several reversible and irreversible modifications. These post-translational modifications (PTMs) result in the diversification of the protein functions beyond what is dictated by gene transcripts. Through these PTMs, eukaryotic cells can dynamically alter the structure, stability and function of a wide spectrum of proteins. Cleavage, addition of sugars, lipids, small proteins or chemical groups such as methyl, nitric oxide, phosphate and acetate (Fig. 3.1), are well known PTMs that regulate the functional expression of ion channels in different physiological and physiopathological con-

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**Fig. 3.1** Schematic representation of some post-translational modifications in ion channels. Addition of complex molecules, small proteins, chemical groups and cleavage is depicted. In this chapter we discuss the role of N-glycosylation and phosphorylation in TRP channels. (Inspired in Fig. 1 of (Wang et al. 2014))



texts. The regulation of ion channels function occurs either modulating their subcellular trafficking or altering their biophysical properties. TRP channels are not the exception, and among the PTMs that exert an enormous effect on their physiological role, N-glycosylation and phosphorylation stand out. In sensory transduction, where several members of this superfamily of ion channels have a critical role, these PTMs regulate their destination to the specialized structures where transduction take place, and tune their contribution to the net response of the sensory cell to different physiologically relevant stimuli.

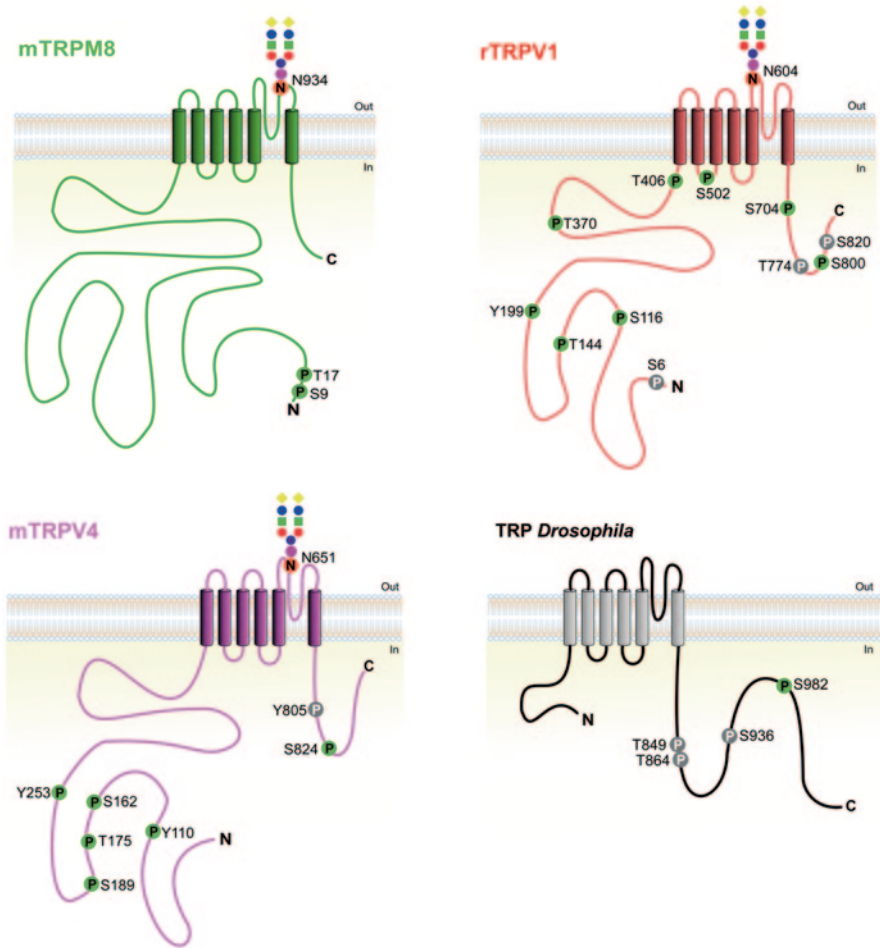
In this chapter, we summarize the contribution of N-glycosylation and phosphorylation to the functional expression of some TRP channels (Fig. 3.2) with a relevant role in sensory transduction.

## 3.2 Modulation of TRP Channels by N-glycosylation

### 3.2.1 The N-linked Glycosylation Process

N-glycosylation is one of the most common PTMs influencing the function of TRP channels. This PTM consists of the covalent addition of sugar residues to asparagines that are part of Asn-X-Ser/Thr consensus sequence, where X represents any amino acid except proline (Apweiler et al. 1999). The sugar composition of this modification results from the activity of different enzymes in the endoplasmic reticulum (ER) and the Golgi apparatus (reviewed by Varki et al. 1999). This process





**Fig. 3.2** Schematic representation of the different monomers of *mouse TRPM8*, *rat TRPV1*, *mouse TRPV4* and *Drosophila TRP*. Phosphorylated residues, identified with specific antibodies, site directed mutagenesis and amino acid sequencing are represented by *P*. Other phosphorylated residues identified by mass spectrometry are included in Table 3.1. *Green* and *grey* backgrounds in *P* represent phosphorylation with or without reported effects in channel function, respectively. N-glycosylation is shown as a branched chain of sugar moieties in asparagine residues (*N*)

begins with the co-translational transference of the oligosaccharide precursor Glc3Man9GlcNAc2 to the nascent polypeptide (reviewed by Varki et al. 1999). The acceptor asparagine has to be accessible to the oligosaccharyltransferase complex, which catalyzes the transference of the oligosaccharide precursor to the polypeptide on the luminal side of the ER membrane. The resulting glycoprotein undergoes enzymatic removal of three terminal glucose moieties in the ER, resulting in the “high-mannose” state, and is then transported to the Golgi apparatus where additional modifications occur (reviewed by Varki et al. 1999). In this compart-



ment, different enzymes may modify the early/immature glycosylation, generating a complex mature N-glycosylation. Unlike the high mannose glycosylation that takes place in the ER, which is the same for all the proteins, the composition of the N-glycosylation tree displayed at the end of the transit of the proteins through the Golgi is quite diverse, and depends on the specificities of Golgi enzymes involved and their tissue-specific expression patterns (Schachter et al. 1983; Comelli et al. 2006).

Several reports have shown that glycans can modify functional properties of ion channels (see (Scott and Panin 2014) for a review). One of the mechanisms by which N-glycosylation regulates ion channels function is modulating their trafficking to the cell surface. The role of this PTM in processes such as protein folding, stability and subcellular trafficking are critical to determine the final density of ion channels in the plasma membrane (reviewed by Helenius and Aebi 2001). N-glycosylation can also alter diverse biophysical properties of the channel, tuning their physiological function. Both modulatory mechanisms can be found in TRP channels (reviewed by Cohen 2006). For instance, the hydrolysis of some sugar residues of TRPV5 N-glycosylation by *klotho*, a glucuronidase enzyme, results in the stabilization of the channel at the apical plasma membrane of kidney epithelial cells, increasing the expression of TRPV5 in this compartment (Chang et al. 2005). This mechanism is used to regulate reabsorption of  $\text{Ca}^{2+}$  in the nephron, that maintains the normal  $\text{Ca}^{2+}$  levels in the blood (Chang et al. 2005). On the other hand, in the case of TRPC6 and TRPC3, differences in N-glycosylation pattern are responsible for basal channel activity (Dietrich et al. 2003). Mutation of one of the two N-glycosylation sites of TRPC6 turns this tightly regulated channel in a constitutively active channel, similar to TRPC3 (Dietrich et al. 2003).

### 3.2.2 *TRPV1 N-glycosylation*

**Transient Receptor Potential Vanilloid 1 (TRPV1)** is a non-selective, calcium-permeable cation channel, activated by capsaicin, heat, voltage and acidic pH expressed in nociceptors (Caterina et al. 1997). This channel is involved in heat thermosensation and has a critical role in inflammatory pain (Caterina et al. 1997; Davis et al. 2000; Zhang et al. 2005) (see Chap. 8 by McNaughton and Tan in this book). The predicted mass of the TRPV1 monomer is ~95 kDa, however, western blot experiments of transfected cell lines reveal the presence of higher molecular weight bands, suggesting that TRPV1 is a N-glycosylated protein (Kedei et al. 2001; Jahnel et al. 2001). Treatment of protein extracts with different glycosidases corroborated the existence of an immature high mannose type glycosylation that takes place in the ER, and a mature (or complex) glycosylation that requires protein transit through the different Golgi compartments and generates a diffuse band at ~115 kDa (Kedei et al. 2001; Jahnel et al. 2001; Veldhuis et al. 2012). Using biotinylation in a recombinant system, the analysis of TRPV1 channels located in the plasma membrane also evidenced the presence of multiple bands, suggesting that the channel at the cell surface could exist in different states of glycosylation (Veldhuis et al. 2012).

While the N-glycosylation of TRPV1 in transfected cell lines, such as **H**uman **E**mbrionic **K**idney (**HEK293**) cells and the dorsal root ganglia derived cell line F11, has been reported by several laboratories (Kedei et al. 2001; Jahnel et al. 2001; Veldhuis et al. 2012), the N-glycosylation of the native TRPV1 is more controversial. This is not a trivial point, since this kind of PTM is strongly dependent on the cellular context and it must be considered in order to select the appropriate expression system to study the functional properties of an ion channel of interest, and especially to extrapolate these findings to a physiological function of native channels. Using western blot analysis, Kedei and co-workers have reported the lack of mature N-glycosylation in purified membrane extracts of dorsal root ganglia (DRG) (Kedei et al. 2001). In contrast, Veldhuis and co-workers showed that N-glycosidase activity affected the apparent molecular weight of TRPV1 from rat DRG and sciatic nerve, suggesting that this post-translational modification occurs *in vivo* (Veldhuis et al. 2012).

The N-glycosylation of TRPV1 channels takes place at residue Asn604, located in the third extracellular loop near the pore domain (Jahnel et al. 2001). Although the lack of this PTM does not affect TRPV1 trafficking to the plasma membrane (Veldhuis et al. 2012), it significantly alters some of its biophysical properties (Wirkner et al. 2005; Veldhuis et al. 2012). Electrophysiological experiments revealed that mutation of N604T, that prevents the N-glycosylation, causes a significant reduction in the  $EC_{50}$  value for capsaicin and in the maximal response of the channel to this agonist (Wirkner et al. 2005). The response to 1  $\mu$ M capsaicin of the unglycosylated channel is much more sensitive to changes in extracellular proton concentration, suggesting that this PTM plays a role in the regulation of the TRPV1 pH sensitivity (Wirkner et al. 2005). Interestingly, compared to the wild type channel, in the N604T mutant the antagonist capsazepine was a more effective blocker of capsaicin-induced responses at physiological pH (Wirkner et al. 2005). Thus, these data correlate the presence of N-glycosylation with a decrease in TRPV1 sensitivity to agonists such as capsaicin and protons, and antagonists like capsazepine. During inflammatory responses, alterations in extracellular pH are common, and acidic pH changes predominate. In these conditions, N-glycosylation of TRPV1 dampens the effect of acidic pH on channel activation by other physiologically relevant stimuli, such as heat or endogenous modulators of its function in nociceptors. Thus, thermal hyperalgesia that results of an inflammatory state could be reduced if the channels are mainly in the N-glycosylated form in nociceptive free nerve endings.

The role of N-glycosylation in the capsaicin-evoked pore dilation, that has been recently described in TRPV1 channels (Chung et al. 2008), has also been investigated. Chung and co-workers described how activation of TRPV1 with capsaicin leads to an increase in relative permeability to large cations and  $Ca^{2+}$  through agonist-induced pore dilation process, a structural change originally described in ATP-gated P2Xs channels, which has also been studied in other TRP channels (Chen et al. 2009). The effect of N-glycosylation of TRPV1 in the pore dilation mechanism was assessed by Veldhuis and co-workers, measuring the capsaicin-dependent uptake of the large fluorescent cationic dye YO-PRO-1. Cells transfected with the unglycosylated mutant showed a decrease in the uptake of this molecule, suggesting that the pore dilation state induced by capsaicin is in part dependent on the N-glycosylation

status of TRPV1 (Veldhuis et al. 2012). Thus, capsaicin-induced pore dilation and the subsequent enhanced permeability to  $\text{Ca}^{2+}$  ions could be critical for the selective death of TRPV1-expressing neurons. Prolonged exposure to this vanilloid is one of the most effective treatments used to reduce pain in several neuropathies, and pore dilation may contribute to the paradoxical analgesic effect of capsaicin though death of nociceptive sensory nerve fibers (Bautista and Julius 2008).

The N-glycosylation of TRPV1 has also been connected with channel desensitization. Analysis of the decay of the responses to sustained stimulation with capsaicin has shown that the desensitization rate of the response in cells expressing TRPV1 displayed a large variability. In contrast, the response of the cells transfected with the unglycosylated N604T mutant always showed a constant relaxation kinetic (Veldhuis et al. 2012). This observation suggests that desensitization depends, at least to some extent, on the glycosylation status of the channel. As mentioned above, TRPV1 located in the plasma membrane shows different states of glycosylation, that could account for the observed cell-to-cell variability (Veldhuis et al. 2012). How this variability may influence the physiological net response of TRPV1 channels in native membranes is still unclear.

### 3.2.3 *TRPM8 N-glycosylation*

**Transient Receptor Potential Melastatine 8 (TRPM8)** is a thermoTRP channel activated by cold, cooling compounds such as menthol and by voltage. This channel is the main molecular entity responsible for detection of cold temperatures in the somatosensory system (recently reviewed by McCoy et al. 2011; Almaraz et al. 2014; Madrid and Pertusa 2014); see also Chap. 9 by González et al. in this Book). Like TRPV1, TRPM8 is also a glycoprotein (Dragoni et al. 2006; Erler et al. 2006; Tsuruda et al. 2006). In TRPM8, the N-glycosylation takes place at residue Asn934, located in the extracellular loop between transmembrane domains five and six, near the pore domain (Dragoni et al. 2006; Erler et al. 2006). Analysis of protein extracts of HEK293 cells transfected with TRPM8 shows three distinguishable bands corresponding to different glycosylation states of TRPM8: the unglycosylated protein, an early N-glycosylated form produced in the ER and the mature N-glycosylated form that contains terminal sialic acid residues (Morenilla-Palao et al. 2009; Pertusa et al. 2012). The N934Q mutant, which has been used to evaluate the effect of this PTM in TRPM8 activity, shows a reduced response to cold and menthol (Dragoni et al. 2006; Erler et al. 2006; Pertusa et al. 2012). Different mechanisms have been proposed to explain this N-glycosylation-dependent effect on TRPM8. Whereas some authors suggest that it is mainly the result of a drop in the number of TRPM8 channels in the plasma membrane (Dragoni et al. 2006; Erler et al. 2006), others have suggested a direct effect on N-glycosylation on the biophysical properties of the channel (Pertusa et al. 2012). It has been demonstrated that the absence of this PTM causes a shift in the voltage-dependence of the channel of almost 60 mV to more positive voltages, decreasing its open probability at physiologically relevant

membrane potentials (Pertusa et al. 2012), therefore influencing its cold and menthol responses.

Protein extracts from DRG and trigeminal ganglia (TG) revealed that TRPM8 N-glycosylation also takes place *in vivo*. In fact, the treatment of TG neurons with the N-glycosylation inhibitor tunicamycin shifts the mean temperature threshold of cold responses from  $\sim 29$  to  $\sim 23$  °C (Pertusa et al. 2012). This shift in recombinant systems is from  $\sim 26$  to  $\sim 23$  °C, suggesting a larger impact of this PTM in temperature-dependent responses in native channels (Pertusa et al. 2012).

N-glycosylation also plays a role in the segregation of TRPM8 to lipid rafts (Morenilla-Palao et al. 2009), facilitating its association to these plasma membrane microdomains that contain high concentrations of cholesterol and glycosphingolipids (Simons and Ikonen 1997). It has been reported that within the plasma membrane TRPM8 is mainly localized in lipid rafts, in both native and recombinant systems (Morenilla-Palao et al. 2009). Interestingly, the association of TRPM8 to lipid rafts decreases both menthol- and cold-mediated activation of the channel. This observation suggests that TRPM8 can be affected by its lipid environment and that menthol- and cold-mediated responses of this channel are facilitated when the protein is outside these domains (Morenilla-Palao et al. 2009). Therefore, unglycosylated channels, which show an important reduction in their association to lipid rafts (Morenilla-Palao et al. 2009), should display higher responses to agonists than the N-glycosylated protein. Nevertheless, as mentioned before, the non-glycosylated mutant shows a reduced response to cold and menthol stimulation, indicating that N-glycosylation *per se*, independent of the lipid raft association, has an important influence on biophysical properties of TRPM8 (Pertusa et al. 2012).

Lipid rafts have been linked to vital cell processes such as trafficking and sorting of membrane proteins (reviewed by Pristera et al. 2012). Although the roles of lipid rafts and N-glycosylation in protein sorting of TRPM8 still remains unexplored, this association may be critical for the correct targeting and trafficking dynamics in native membranes. Moreover, it is also important to consider that by promoting TRPM8 association with lipid rafts, N-glycosylation may also be favoring protein-protein interactions that could modulate channel function in the nerve endings of primary somatosensory neurons, where cold transduction occurs. Thus, N-glycosylation of TRPM8 emerges as an important mechanism that can modulate channel function, affecting its biophysical properties and determining the association of the channel with lipid rafts.

Altogether, these results illustrate the relevance of N-glycosylation in the physiological function of TRPM8 and suggest that in native systems it could be critical in the fine tuning of cold sensitivity in cold thermoreceptor neurons.

### 3.2.4 TRPV4 N-glycosylation

As TRPM8 and TRPV1, **Transient Receptor Potential Vanilloid 4 (TRPV4)** is also a polymodal TRP channel activated by a plethora of stimuli, including cell swelling,

innocuous heat, arachidonic acid, the endogenous cannabinoid neurotransmitter anandamide and several exogenous ligands. TRPV4 is a broadly expressed channel with important roles in osmosensation and thermosensation, and mutations of the gene encoding this protein are related to diverse diseases (see Nilius and Voets 2013 for a complete review).

As other glycosylated TRP channels, immunoblots from heterologously expressed and native TRPV4 show multiple bands (Xu et al. 2006; Arniges et al. 2006). In agreement with the presence of N-glycosylation, treatment of cell lysates with endoglycosidase F results in a single band at 96 kDa, corresponding to the non-glycosylated form of the channel (Xu et al. 2003; Arniges et al. 2006). Like in TRPV1 and TRPM8, western blots of TRPV4 show two additional higher molecular weight bands with apparent molecular masses of 100 and 110 kDa, representing respectively the high mannose and complex glycosylation states of the channel (Arniges et al. 2006). Site-directed mutagenesis of potential glycosylation sites revealed that N-glycosylation of TRPV4 occurs at residue Asn651, located in the pore loop (Xu et al. 2006). The unglycosylated mutant channel N651Q, expressed in HEK293 cells, exhibited greater responsiveness to hypotonicity (Xu et al. 2006). The mechanism behind this effect was revealed using biotinylation assays, where a higher expression level of this mutant channel at the cell surface was detected. Therefore, unlike the previous examples in the other thermoTRP channels mentioned above, the alteration in the response observed in the unglycosylated mutant is most likely due to enhanced trafficking of the protein to the plasma membrane.

Thus, N-glycosylation emerges as a critical PTM influencing the physiological role of TRP channels that are key elements in thermotransduction, inflammatory pain and transduction of osmotic stimuli.

### 3.3 Regulation of TRP Channels by Phosphorylation

#### 3.3.1 Protein Phosphorylation

Protein phosphorylation consists in the transferring of a phosphate to the hydroxyl group of a serine, threonine or tyrosine of a substrate protein, usually producing a modification in its functional properties. This reversible and covalent modification is catalyzed by different kinases, and dephosphorylation is carried out by phosphoprotein phosphatases. Ion channels are targets of these enzymes, and the addition or removal of phosphate groups can modify the channel activity determining the net electrophysiological properties of the cell (reviewed by Swope et al. 1992). Typically, the phosphorylation pattern of ion channels involves several sites with consensus sequences for different protein kinases. TRP channels are modulated by phosphorylation, with quite diverse consequences. For example, TRPC3 and TRPC6 are inhibited by PKC-mediated phosphorylation, in contrast with the enhancement observed after Src family tyrosine kinases (SFK)-dependent phosphorylation (reviewed by Woolstra and Huber 2014).

In the next section we will discuss some examples of modulation by phosphorylation in TRPs involved in sensory transduction, with especial attention to TRPV1 channel, whose modulation by this PTM has been extensively studied.

### 3.3.2 *Phosphorylation in TRPV1*

#### 3.3.2.1 **Role of Phosphorylation in TRPV1 Channel Sensitization**

Most of the nociceptive neurons show an increase in their responses under repetitive noxious stimulation. In these neurons, TRPV1 plays a key role in the generation of hyperalgesia, defined as exacerbated pain in response to normally painful stimulation, occurring under pathological states such as inflammation (Caterina et al. 1997; Davis et al. 2000). TRPV1 is directly responsible for the hyperalgesia triggered by several proinflammatory mediators including bradykinin, nerve growth factor (NGF) and ATP (Caterina et al. 1997; Cesare et al. 1999; Shu and Mendell 1999; Premkumar and Ahern 2000; Chuang et al. 2001; Vellani et al. 2001; Prescott and Julius 2003; Bonnington and McNaughton 2003). The interaction of these proinflammatory factors with their receptors leads a downstream activation of different kinases, such as PKC, PKA or SFKs. These kinases phosphorylate specific residues in TRPV1, causing an enhanced response of this channel to heat, capsaicin, protons or anandamide (Premkumar and Ahern 2000; Vellani et al. 2001; Numazaki et al. 2002; Zhang et al. 2005). Thus, phosphorylation of TRPV1 enhances its responses to endogenous and exogenous agonists, either by modifying its biophysical properties or by increasing the number of channels in the plasma membrane (Zhang et al. 2005, 2008; Studer and McNaughton 2010).

PKC is essential to the enhanced response of TRPV1 during inflammatory hyperalgesia (Premkumar and Ahern 2000; Vellani et al. 2001). Among the different putative targets of PKC in the protein, the phosphorylation of Ser502 and Ser800 are critical for this effect (Numazaki et al. 2002; Bhave et al. 2003). Mutation of Ser502 and Ser800 to alanine prevents both channel phosphorylation and the potentiation of the TRPV1-dependent responses caused by the PKC activator phorbol 12-myristate 13-acetate (PMA) (Numazaki et al. 2002; Bhave et al. 2003). Electrophysiological evidence obtained at single-channel level of the double mutant S502A/S801A of human TRPV1 has been useful to understand the mechanism behind this potentiation (Studer and McNaughton 2010). Activation of PKC enhances the response of TRPV1 to capsaicin, increasing the open probability of the channel with no alterations in single-channel conductance (Studer and McNaughton 2010). One possible explanation for this effect is that Ser502/Ser801 phosphorylation increases the capsaicin binding affinity. However, phosphorylation of TRPV1 by PKC also potentiates the response to heat, protons and anandamide (Premkumar and Ahern 2000; Vellani et al. 2001) The intracellular capsaicin binding site is also implicated in anandamide activation (Jordt and Julius 2002), while protons target an



extracellular site (Jordt et al. 2000). Given the distance between these two activation sites, it seems unlikely that the increased TRPV1 activity following PKC phosphorylation is only due to changes in the conformation of the capsaicin binding site. This suggests that phosphorylation of these residues affects common downstream steps that lead to channel opening (Studer and McNaughton 2010). On the other hand, it is important to point out that in basal conditions the double mutant S502A/S801A and the wild type channel are similarly activated by capsaicin. This suggests that the phosphorylation of these two serines are only involved in the potentiating effect of PKC (Studer and McNaughton 2010). In addition, mentioned before, agonist stimulation induces dynamic changes in the ionic selectivity of TRPV1 that could be enhanced by phosphorylation of Ser800 by PKC (Chung et al. 2008). This mechanism may contribute to the modulatory effects of different inflammatory mediators on TRPV1 function in nociceptors (Chung et al. 2008).

Nevertheless, other mechanisms could also be involved in the potentiation of PKC stimulation of TRPV1-dependent responses. TRPV1 channels located in vesicles can be rapidly recruited to the plasma membrane by regulated exocytosis upon PKC stimulation, increasing their density at the plasma membrane (Morenilla-Palao et al. 2004). In fact, it has been proposed that the interaction of the channel with AKAP79/150 (see below), together with the phosphorylation of TRPV1 in Ser502 by PKC or PKA, is required to promote TRPV1 trafficking to the cell surface (Zhang et al. 2008).

NGF also causes an enhanced trafficking of TRPV1 to the plasma membrane. The interaction of NGF with the TrkA receptor initiate a signalling pathway that activates the Src kinase, which directly phosphorylates TRPV1 at Tyr200, promoting the insertion of channel in the plasma membrane (Zhang et al. 2005). As a consequence, the responses of TRPV1 to agonists after NGF exposure are larger, influencing the net response of the nociceptive neuron to endogenous and exogenous activators of TRPV1 in hyperalgesic events (Zhang et al. 2005).

### 3.3.2.2 Basal Phosphorylation of TRPV1 and Desensitization

Phosphorylation of TRPV1 can also occur in basal conditions. In fact, several studies have reported that TRPV1 transfected cells display basal  $^{32}\text{P}$  incorporation, indicating that this channel is constitutively phosphorylated (Numazaki et al. 2002; Bhave et al. 2002; Jung et al. 2004; Jeske et al. 2006). In agreement with these results, the general kinase inhibitor staurosporine reduces single-channel activity, supporting the relevance of basal phosphorylation on TRPV1 activity (Studer and McNaughton 2010).

One remarkable feature of TRPV1 is the pronounced desensitization during sustained agonist exposure (acute desensitization), or upon repetitive agonist applications (tachyphylaxis). Interestingly, both effects are correlated with a robust dephosphorylation of the protein after activation (Docherty et al. 1996; Bhave et al. 2002). The desensitization of capsaicin-evoked currents is dependent on external  $\text{Ca}^{2+}$  (Docherty et al. 1996; Koplas et al. 1997) and requires  $\text{Ca}^{2+}$ - and calmodulin-



dependent protein phosphatase 2B (calcineurin) (Docherty et al. 1996; Jung et al. 2004; Mohapatra and Nau 2005). Among the residues constitutively phosphorylated involved in the desensitization, Ser116 was the first identified (Bhave et al. 2002). Bhave and colleagues demonstrated that this position is dephosphorylated after capsaicin exposure, leading TRPV1 desensitization. Tachyphylaxis could be prevented by PKA activation that re-phosphorylates Ser116 (Bhave et al. 2002). However, dephosphorylation and phosphorylation of Ser 116 do not represent desensitized or sensitized states of the channel. In contrast, a mechanism was suggested where a dephosphorylated Ser116 allows the entry of TRPV1 into a desensitized state, which is prevented by the phosphorylation of the channel in this position (Bhave et al. 2002).

PKA also phosphorylate Thr370 and Thr144 in TRPV1 (Bhave et al. 2002). Interestingly, mutants T370D, T370A, T144D T144A exhibited a significantly reduced acute desensitization and tachyphylaxis. However, treatments with the PKA activator forskolin, or calcineurin inhibitors, used to reduce these desensitization processes, had different effects in these mutants. In the Thr144 mutants, both strategies caused a decrease in tachyphylaxis, whereas in Thr370 mutants, these treatments were ineffective. These results suggest that only Thr370 is involved in the reduction of the desensitization induced by PKA activation (Mohapatra and Nau 2003, 2005). The phosphorylation of residues Thr370 and Thr144 of TRPV1 has also been related with the anti-hyperalgesic effect displayed by some cannabinoid agonists such as WIN 55,212-2 (WIN). These residues are constitutively phosphorylated in native TRPV1 from DRG neurons (Jeske et al. 2006). In the subset of somatosensory neurons that coexpress TRPV1 and TRPA1 channels, activation of TRPA1 by WIN causes a  $Ca^{2+}$  influx that activates calcineurin, leading to the dephosphorylation of these residues and desensitization of TRPV1 (Jeske et al. 2006).

It has also been reported that native TRPV1 can be directly phosphorylated at position Thr407 by cyclin-dependent kinase 5 (**cdk5**) (Pareek et al. 2007). When this kinase is inhibited, capsaicin responses of the channel are significantly smaller (Pareek et al. 2007). Thus, cdk5 in DRG neurons would be responsible for a basal phosphorylation of TRPV1 that increase its sensitivity to agonists in a native system (Pareek et al. 2007).

Finally, it has been described that Thr705 is phosphorylated in the subset of DRG neurons that coexpress TRPV1 and PKC $\beta$ II. The direct binding of the PKC $\beta$ II to the channel activates this kinase, which in turn phosphorylates Thr705 producing an increase in the sensitivity of TRPV1 to heat and capsaicin. Thus, in basal conditions, DRG neurons display different temperature thresholds depending on whether TRPV1 is expressed simultaneously with PKC $\beta$ II or not (Li et al. 2014).

In addition, dephosphorylation of Thr704 has been related with TRPV1 desensitization. Experiments with the double mutant S502A/T704I have suggested that the dephosphorylation of both positions causes a reduction in capsaicin binding (Jung et al. 2004).

Altogether, these results suggest that in a non-stimulated cell, TRPV1 displays a basal phosphorylation that depends on cellular context and strongly determines the functional properties of the channel in basal conditions, influencing its physiological role in native membranes.

### 3.3.2.3 TRPV1 Phosphorylation and Scaffolding Proteins

The speed and specificity of the action of kinases is favoured in many cases by scaffolding proteins, which assemble these enzymes into signalling complexes with their substrate proteins. The scaffolding protein AKAP (A Kinase Anchoring Protein) integrate a large number of kinases and phosphatases into functional signalling complexes (Reviewed by (Smith et al. 2006)). AKAP79/150 has three orthologs: human AKAP79, rodent AKAP150 and bovine AKAP75. AKAP79/150 links PKA, PKC and calcineurin to different ion channels, including glutamate receptors, calcium channels, and M-type potassium channels (Gao et al. 1997; Colledge et al. 2000; Altier et al. 2002; Hoshi et al. 2003; Sandoz et al. 2006; Oliveria et al. 2007; Chai et al. 2007). Early works suggested that an AKAP protein may be involved in the modulation of TRPV1 by PKA (Rathee et al. 2002; Distler et al. 2003), and more recent evidence demonstrated a direct interaction of TRPV1 and AKAP79/150 in native systems (Zhang et al. 2008; Schnizler et al. 2008; Jeske et al. 2008).

Zhang and co-workers showed that overexpression of AKAP79 enhanced the sensitivity of TRPV1 to capsaicin. AKAP deletions without PKC binding site shows an important diminution in the responses to capsaicin, suggesting that PKC phosphorylation is important to modulate TRPV1 in basal conditions. However, a deleted version of AKAP79 protein lacking the PKA binding site has a smaller effect, suggesting that phosphorylation by PKA is less important than that of PKC at resting conditions (Zhang et al. 2008). AKAP79 also participates in the sensitization of TRPV1 by proinflammatory mediators that modulate the channel via PKA and PKC (Zhang et al. 2008; Schnizler et al. 2008; Jeske et al. 2008). In fact, the simple activation of these kinases is not enough to cause potentiation of TRPV1; the generation of a scaffolding complex between TRPV1, AKAP79, and PKC (or PKA) is required for sensitization (Zhang et al. 2008). In addition, AKAP79/150 is also implicated in TRPV1 desensitization through the binding site for the calcium-dependent phosphatase calcineurin (Zhang et al. 2008).

Summarizing, phosphorylation of the heat- and capsaicin-activated TRPV1 channel plays a critical role in fine tuning its response to thermal and chemical stimulation in basal conditions, or mediating the reversible regulation that occurs during sensitization triggered by proinflammatory mediators in nociceptive neurons.

### 3.3.3 Phosphorylation in TRPM8

In the case of the cold- and menthol-activated channel TRPM8, its function can be modulated by different protein kinases. In contrast to TRPV1, the activation of PKC by bradykinin or PMA produces a downregulation of TRPM8 (Premkumar et al. 2005; Abe et al. 2006; Linte et al. 2007), together with a decrease in serine-phosphorylation (Premkumar et al. 2005), suggesting that in basal conditions is phosphorylated. However the kinases that participate in this constitutive

phosphorylation, the residues where it takes place and its physiological relevance remain unresolved.

Different studies have also correlated a decrease in TRPM8 phosphorylation with a reduction in its sensitivity to chemical stimulation. Bavencoffe et al. (2010) reported a PKA-dependent basal phosphorylation of Ser9 and Thr17 residues of TRPM8. In this work, activation of the G<sub>i</sub> protein and subsequent inhibition of the adenylate cyclase (AC)/cAMP/PKA pathway results in a decrease in the PKA-dependent phosphorylation, causing a reduction of the channel responses (Bavencoffe et al. 2010). However, the effects of G protein-PKA activation on TRPM8 function are diverse. Other authors have reported the opposite effect: activators of the AC/cAMP/PKA pathway, such as forskolin and 8-Br-cAMP, reduced the responses to agonists of the channel (De Petrocellis et al. 2007; Linte et al. 2007). Moreover, PKA is involved in TRPM8 inhibition by prostaglandin E2 (Linte et al. 2007).

Thus, although several reports suggest a relevant modulation of TRPM8 by phosphorylation, the precise mechanism behind this regulation is still poorly understood.

### 3.3.4 Phosphorylation in TRPV4

The osmo- and thermo-sensitive channel TRPV4 is also modulated by phosphorylation. PKC and PKA direct phosphorylation of TRPV4 causes a potentiation of the responses to hypotonic stimuli (Fan et al. 2009). The residues involved in the enhanced responses of the channel induced by PKC activation are Ser162, Thr175 and Ser189 in the N-terminal domain, and Ser824 in the C-terminal domain. The phosphorylation of the latter position seems to be involved not only in PKC but also in PKA-dependent potentiation (Fan et al. 2009; Peng et al. 2010). Substitution of Ser824 by an aspartic acid, mimicking a phosphorylation in this site, increases TRPV4-mediated calcium influx in basal conditions and under hyposmotic stimulation, stressing the relevance of this residue in TRPV4 regulation (Peng et al. 2010). Like in TRPV1, the scaffolding protein AKAP79 recruits PKC, PKA and TRPV4, facilitating the phosphorylation of the channel required to channel sensitization (Fan et al. 2009). In addition, TRPV4 is also modulated by SFKs (Xu et al. 2003; Wegierski et al. 2009). An early study showed that hypotonic stress results in a rapid tyrosine phosphorylation of TRPV4 channel in both native and heterologous systems (Xu et al. 2003). In this work, it was suggested that this phosphorylation is carried out by Lyn, a kinase activated by hypotonic stress (Xu et al. 2003). The site of tonicity-dependent tyrosine phosphorylation was identified by mutagenesis as Tyr253 (Xu et al. 2003). The Y253F mutant lost the hypotonicity-dependent phosphorylation and the TRPV4 response to hypotonic stimuli (Xu et al. 2003). However, other evidence has shown that this mutant displays similar responses to hypotonic stress and to other TRPV4-activating stimuli than in the wild type, questioning the role of Tyr253 phosphorylation in TRPV4 activation by hypotonic stress (Vriens et al. 2004).

Other studies support a relevant role of SFKs in the regulation of TRPV4. Two phosphorylation sites in the cytosolic N- and C-terminal domains of this channel were identified by mass spectrometry, using TRPV4 purified from HEK293 and MCDK at normal cell culture conditions. The analysis showed that in unstimulated cells, tyrosines Tyr110 and Tyr805 are phosphorylated by endogenous SFKs (Wegierski et al. 2009). However, calcium imaging experiments using Y110F and Y805F mutants, revealed that only phosphorylation of Tyr110 would be involved in the modulation of the responses of TRPV4 to heat, mechanical stress, and hypotonic cell swelling (Wegierski et al. 2009). As in TRPV1, direct phosphorylation of TRPV4 by PKC, PKA and SFKs produces sensitization of the channel, that could be relevant in inflammatory hyperalgesia (Fan et al. 2009; Wegierski et al. 2009). Thus, understanding how the phosphorylation pathways affect TRPV4 channel function is critical to clarify the role of this polymodal channel in pathological pain conditions.

### 3.3.5 Phosphorylation in *Drosophila* TRP

The founding member of the superfamily of TRP channels, the *Drosophila* TRP channel (see Chap. 5, by Bacigalupo et al. in this book), is expressed in the photosensitive rhabdomeral membrane of photoreceptor cells. This channel is the target of the phospholipase C-mediated visual transduction cascade, together with a second channel, termed TRPL (TRP-like) (reviewed by Minke and Parnas 2006). Light-activated rhodopsin couples to an heterotrimeric G protein, activating phospholipase C $\beta$  (PLC), which hydrolyses PIP<sub>2</sub> (phosphatidylinositol 4,5-bisphosphate) generating inositol 1,3,5-triphosphate (IP<sub>3</sub>), 1,2-diacylglycerol (DAG) and a proton. The production of DAG also activates the eye-specific protein kinase C (eye-PKC), essential for deactivating the light response (reviewed by Minke and Parnas 2006). *Drosophila* visual signalling is one of the fastest transduction cascades (Zuker 1996), partly due to the formation of a macromolecular complex that increases the efficiency of the signal transduction. This complex is organized by the scaffolding protein INAD that holds together the TRP channel, PLC and an eye-specific PKC (Huber et al. 1996). This protein complex is also required for the rapid termination of the light response (Popescu et al. 2006). In fact, the fast component of the deactivation process depends on the direct phosphorylation of TRP channel at Ser982 by eye-PKC, facilitated by the interaction of both proteins with INAD (Popescu et al. 2006). However comparison of transgenic flies expressing a modified *trp*, *trp*<sup>S982A</sup> or a mutant lacking eye-PKC, revealed that the deactivation defect in the latter is more complex than displayed by *trp*<sup>S982A</sup>, which only became evident upon application of very bright light stimulus. This suggests that phosphorylation of additional PKC sites in TRP or other targets may also contribute to the fast deactivation of the visual response (Popescu et al. 2006).

To identify other TRP phosphorylation sites that can also modulate the light response, a recent study used a quantitative mass spectrometry approach where samples from light- and dark- adapted flies were analyzed (Voolstra et al. 2010,

2013). Twenty-eight phosphorylation sites were identified, one serine (S15) in the N-terminus and 27 other sites within the C-terminal domain (Voolstra et al. 2010, 2013). Among the C-terminal phosphorylation sites, 15 displayed enhanced phosphorylation light conditions, whereas position Ser936 exhibited an increase in the phosphorylation in the dark (Voolstra et al. 2010). Among the positions presenting an enhanced phosphorylation state with light, Thr849 and Thr864 stand out (Voolstra et al. 2013). Experiments with different mutants of the phototransduction cascade that exhibit impaired vision show that this phosphorylation is triggered by the calcium entry through TRP channels. Experiments using a constitutively active mutant of TRP channel showed an increased phosphorylation even in dark conditions. However, mutants with a completely functional phototransduction cascade but expressing a non-conducting TRP variant show total lack of phosphorylation, regardless of the light conditions (Voolstra et al. 2013). Despite the identification of these phosphorylation sites of the *Drosophila* TRP channel, more studies are needed to fully understand the role of these post-translational modifications in vivo.

### 3.4 Concluding Remarks

In summary, multiple lines of evidence show that several TRP channels are regulated N-glycosylation and phosphorylation, and that these PTMs are part of the molecular determinants involved in setting their function. N-glycosylation and basal phosphorylation modulate their responses to agonists, tuning their activity or directing these channels to specific membrane compartments in different cells. Inducible phosphorylation, in contrast, works as a dynamic and reversible mechanism responsible for modifications of TRP channel function, a modulation that could be relevant in pathological states such as inflammatory pain. Although our knowledge on the role of these PTMs in TRP channel function has progressed significantly, there are several questions that remain unclear. For example, some residues where phosphorylation actually occurs have been identified, but their contribution to channel function remains elusive (Table 3.1). There is also evidence suggesting that these channels are modulated by a kinase/phosphatase activity, but the position/s of putative phosphorylations have not been identified. More efforts are needed to fully understand the real relevance of these modulations in vivo under physiological and physiopathological conditions.

**Acknowledgments** Supported by Grants FONDECYT 11130144 (MP), FONDECYT 1131064 (RM) and CONICYT Anillo ACT-1113 (MP, RM).

**Table 3.1** Phosphorylated and N-glycosylated residues identified in depicted TRP channels

<i>Residue</i>	<i>Species</i>	<i>Method</i>	<i>Effect</i>	<i>Reference</i>
<i>TRPV1</i>				
Ser6 (P)	r	S	NRE	(Bhave et al. 2002)
Ser116 (P)	r	S	Dephosphorylation of Ser 116 allows desensitization of TRPV1 response	(Bhave et al. 2002; Mohapatra and Nau 2003, 2005)
Thr144 (P)	r	S	Minor phosphorylation site. T144A mutation causes a decreased desensitization of TRPV1	(Bhave et al. 2002; Mohapatra and Nau 2003, 2005; Jeske et al. 2006)
Tyr200 (P)	h	S	The phosphorylation of Tyr200, induced by NFG, increased the plasma membrane expression of TRPV1	(Zhang et al. 2005)
Tyr310 (P)	h	MS	NC	(Rikova et al. 2007)
Thr370 (P)	r	S	Thr370 Minor phosphorylation site. Dephosphorylation of Thr370 is involved in the desensitization of TRPV1 response	(Bhave et al. 2002; Mohapatra and Nau 2003, 2005; Jeske et al. 2006)
Thr407 (P)	m	S	Phosphorylation of Thr407 by Cdk5 in DRG neurons, increases capsaicin responses	(Pareek et al. 2007)
Ser502 (P)	h,r	S	Phosphorylation of Ser502 and interaction with the scaffolding protein AKAP79/150 promotes trafficking to the plasma membrane. Phosphorylation of Ser502 by PKC causes an enhanced response of TRPV1 to agonists	(Zhang et al. 2008)
Asn604 (N)	r	S	N-glycosylation affects some biophysical properties of the channel	(Numazaki et al. 2002; Bhave et al. 2003; Studer and McNaughton 2010)
Thr 705 (P)	h	S	PKC $\beta$ II phosphorylates TRPV1 in Thr7045 causing an increase in its temperature sensitivity	(Wirkner et al. 2005; Veldhuis et al. 2012)
Thr704 (P)	r	S	Basal phosphorylation of this residue is required to capsaicin response	(Li et al. 2014)
Thr773 (P)	h	MS	NC	(Jung et al. 2004)
Thr 774 (P)	r	S	NRE	(Hornbeck et al. 2004; Kettenbach et al. 2011)
				(Bhave et al. 2002, 2003)

Table 3.1 (continued)

<i>Residue</i>	<i>Species</i>	<i>Method</i>	<i>Effect</i>	<i>Reference</i>
Thr 775 (P)	h	MS	NC	(Kettenbach et al. 2011)
Ser780 (P)	h	MS	NC	(Kettenbach et al. 2011)
Ser800 (P)	r	S	Phosphorylation of Ser800 by PKC causes an enhanced response of TRPV1 to agonists	(Numazaki et al. 2002; Bhavé et al. 2003)
Ser801 (P)	h	S	Phosphorylation of Ser801 by PKC causes an enhanced response of TRPV1 to agonists	(Studer and McNaughton 2010)
Ser 820 (P)	r	S	NRE	(Bhavé et al. 2002, 2003)
<i>TRPM8</i>				
Ser17 (P)	h	S	Phosphorylation of Ser17 by PKA causes an increase in TRPM8 response	(Bavencoffé et al. 2010)
Asn934 (N)	m,h	S	Mutation of N934 causes a reduction in the responses of TRPM8 to cold and menthol	(Dragoni et al. 2006; Ertler et al. 2006; Pertusa et al. 2012)
<i>TRPV4</i>				
Tyr91 (P)	h	MS	NC	(Hornbeck et al. 2004; Rikova et al. 2007)
Tyr110 (P)	m	S	Phosphorylation in Tyr 110 is involved in modulation of TRPV4 responses to heat, mechanical stress and hypotonic swelling	(Węsierski et al. 2009)
Tyr113 (P)	h	MS	NC	(Hornbeck et al. 2004; Rikova et al. 2007)
Ser162 (P)	h	S	Phosphorylation of Ser162 is involved in the enhanced responses of the channel induced by PKC activation	(Fan et al. 2009)
Thr175 (P)	h	S	Phosphorylation of Ser175 is involved in the enhanced responses of the channel induced by PKC activation	(Fan et al. 2009)
Thr181 (P)	m	MS	NC	(Huttlin et al. 2010)
Ser189 (P)	h	S	Phosphorylation of Ser189 is involved in the enhanced responses of the channel induced by PKC activation	(Fan et al. 2009)
Tyr253 (P)	m	S	Phosphorylation of Tyr253 mediates TRPV4 response to hypotonic stress	(Xu et al. 2003)



Table 3.1 (continued)

<i>Residue</i>	<i>Species</i>	<i>Method</i>	<i>Effect</i>	<i>Reference</i>
Ser505 (P)	h	MS	NC	(Beausoleil et al. 2006)
Tyr508 (P)	h	MS	NC	(Beausoleil et al. 2006)
Asn651 (N)	m	S	Mutation of N-glycosylation sites produces an increase in the expression of TRPV4 in the plasma membrane	(Xu et al. 2006)
Tyr805 (P)	m	S	NRE	(Wegierski et al. 2009)
Ser823	m	MS	NC	(Zamivan et al. 2008)
Ser824	h,m	S	Phosphorylation of Ser824 is involved in PKA and PKC-dependent potentiation	(Fan et al. 2009; Peng et al. 2010)
<i>Drosophila TRP</i>				
Ser15	d	MS	LIP, NC	(Voolstra et al. 2010, 2013)
Ser717	d	MS	NC	(Voolstra et al. 2010, 2013)
Ser721	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)
Ser726	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)
Ser828	d	MS	NC	(Voolstra et al. 2010, 2013)
Thr849	d	S	LDP. The relevance of this phosphorylation has to be determined	(Voolstra et al. 2013)
Thr864	d	S	LDP. The relevance of this phosphorylation has to be determined	(Voolstra et al. 2013)
Ser867	d	MS	NC	(Voolstra et al. 2013)
Ser872	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)
Ser875	d	MS	NC	(Voolstra et al. 2010, 2013)
Ser876	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)
Ser881	d	MS	LIP, NC	(Voolstra et al. 2010, 2013)
Ser884	d	MS	LIP, NC	(Voolstra et al. 2010, 2013)
Ser936	d	S	Phosphorylation of Ser936 occurs in dark conditions. The relevance of this phosphorylation has to be determined	(Voolstra et al. 2010)
Ser956	d	MS	NC	(Voolstra et al. 2010, 2013)
Ser958	d	MS	LDP, NC	(Voolstra et al. 2013)

Table 3.1 (continued)

Residue	Species	Method	Effect	Reference
Ser961	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)
Thr963	d	MS	LIP, NC	(Voolstra et al. 2010, 2013)
Ser964	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)
Ser982	d	S	Phosphorylation of Ser982 is required to the rapid termination of the light response	(Popescu et al. 2006)
Ser990	d	MS	LIP, NC	(Voolstra et al. 2010, 2013)
Thr998	d	MS	LDP, NC	(Voolstra et al. 2013)
Thr1036	d	MS	LDP, NC	(Voolstra et al. 2013)
Thr1049	d	MS	NC	(Voolstra et al. 2013)
Ser1056	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)
Ser1123	d	MS	LDP, NC	(Voolstra et al. 2013)
Ser1253	d	MS	NC	(Voolstra et al. 2013)
Ser1254	d	MS	LDP, NC	(Voolstra et al. 2010, 2013)

MS phosphorylation determined using only a mass spectrometry approach, Specific phosphorylation determined by site directed mutagenesis, specific antibody, and amino acid sequencing, (P) Phosphorylation, (N) N-Glycosylation, h human, r rat, m mouse, d *Drosophila*, NC No characterized, NRE No reported effect, LIP Light independent phosphorylation, LDP Light dependent phosphorylation

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# Chapter 4

## TRP Channels in Visual Transduction

Juan Bacigalupo, Ricardo Delgado, Yorka Muñoz and Peter O'Day

**Abstract** The original members of the superfamily of transient receptor potential channels are the TRP and TRPL channels underlying *Drosophila* light transduction. Light transduction takes place in the photosensitive microvilli within the photoreceptor cells; it is initiated by photon absorption which leads ultimately to the generation of a depolarizing receptor potential caused by TRP and TRPL channel opening. Channel opening is mediated by a phospholipase C pathway where the membrane second messenger diacylglycerol appears to be the channel activator. TRP is a Ca<sup>2+</sup> selective channel responsible for nearly 95% of the net transduction current, whereas TRPL, a poorly-selective Ca<sup>2+</sup> channel, accounts for the difference. The scaffolding protein INAD forms a complex with TRP and other transduction proteins, offering an extremely fast transduction mechanism. TRP and TRPL are also found in the synaptic terminals of the photoreceptors, where they play a role in presynaptic Ca<sup>2+</sup> increments during synaptic transmission.

**Keywords** Light · Transduction · Vision · Retina · Photoreceptor · Lipids · TRP channels

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© Springer International Publishing Switzerland 2015  
R. Madrid, J. Bacigalupo (eds.), *TRP Channels in Sensory Transduction*,  
DOI 10.1007/978-3-319-18705-1\_4

## 4.1 Introduction

The ability of animals to see is remarkable. Even the early steps in vision that involve light detection by photoreceptor cells are very complex and not fully understood in invertebrates and vertebrates. Light detection begins with absorption of photons by photopigments embedded in specialized membranes, and this event triggers biochemical signaling cascades. The nature of these signaling cascades varies widely across species. The biochemical cascades generate transmembrane neural electrical signals that travel to the photoreceptor cell synapse, where the neural information is relayed to the next level of signaling in the visual pathways. The process by which light is coded into an electrical neural signal is called phototransduction, because light energy is transduced into electrochemical energy.

Photoreceptor cell organization and complexity also varies widely, with photoreceptors organized as functional units of simple multicellular tissues or as components of complex eyes that focus light directly to the transducing regions. Some photoreceptors exist as isolated individual cells only serving to detect changes in illumination, forming part of simple photosensitive multicellular arrangements with different degrees of complexity; others are integrated into highly sophisticated organs, such as the compound eyes of insects and the camera eyes of cephalopods and vertebrates. Depending on the photosensitive structure, the animal can distinguish shadows, shapes, colors and even images of great complexity. Visible light represents a small fraction, 400–700 nm, of the vast spectrum of electromagnetic waves that reach our planet,  $10^{-11}$ – $10^3$  cm!

Photoreceptors emerged late in evolution relative to more primitive sensory receptors, such as chemo and mechanotransducing cells. The fact that the number of species boosted dramatically after phototransduction evolved hundreds of million years ago clearly suggests a considerable adaptive value of vision (Arendt 2008; Arendt and Wittbrodt 2001; Lamb 2011).

Photoreceptors are extremely sensitive and remarkably efficient photon detectors. In fact, they can generate discernable electrical responses to single photons and transmit single photon responses to secondary neurons, eventually reaching the central processing regions of the nervous system.

In this chapter we will briefly discuss vertebrate photoreceptors and then shift the focus to *Drosophila melanogaster* photoreceptors, the best characterized of all invertebrate visual cells, which employ TRP channels in phototransduction. Notably, these are the founding members of the extensive TRP channel superfamily, with key roles in a wide variety of sensory transduction mechanisms in invertebrates and vertebrates, such as mechano, thermo, taste, olfactory, pain and osmotransduction.

## 4.2 Vertebrate Photoreceptors

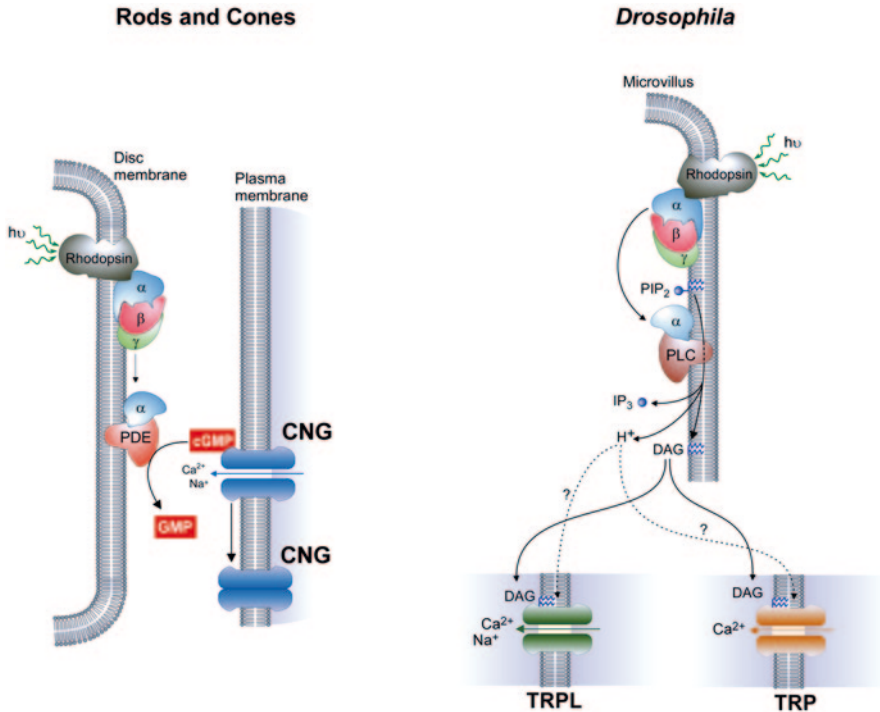
Vertebrate photoreceptors, rods and cones, are widely conserved in function, morphology and biochemical signaling components across species. They are exquisitely arranged and aligned in outer layers of the retina, and they transmit information

in the form of neural signals to higher order cells in more proximal layers. Each cell possesses a clearly distinguishable light transducing region, called the outer segment. This region contains about 2000 internal membrane structures (discs) containing the photopigment rhodopsin, a G-protein coupled membrane receptor, and several other transduction proteins that form the metabotropic signaling cascades that generate and modulate the electrical responses to light. These light responses are mediated by the closing of plasma membrane ion channels, often called transduction channels. These cationic non-selective channels reside exclusively in the outer segment plasma membrane, allowing  $\text{Ca}^{2+}$  and  $\text{Na}^{+}$  ions to enter the outer segment. These are the only ion channel types present in the outer segment. Because these channels are open in the dark, they sustain an inward current in the outer segment that keeps the membrane depolarized. This current is counterbalanced by an outward current at the inner segment membrane through  $\text{K}^{+}$  channels responsible of the receptor potential in most cell types and the membrane potential in darkness is set around  $-35$  mV. As a consequence, there is an extracellular current flow from the inner to the outer segment. Light causes the closure of the transduction channels and the hyperpolarization of the cell due to the reduction of the inward current, shifting the membrane potential towards the  $\text{K}^{+}$  equilibrium potential, reaching nearly  $-70$  mV if challenged with a saturating bright light (Lou et al. 2008).

These transduction channels are gated by cyclic GMP and belong to the small family of the cyclic nucleotide-gated channels, CNGs. cGMP levels are governed by the balance between cGMP production by guanylate cyclase (GC) and cGMP degradation by phosphodiesterase (PDE) that hydrolyzes cGMP to GMP. This balance maintains stable cGMP levels in the dark within the outer segment. Upon absorbing a photon, rhodopsin couples to a GTP-binding protein (G-protein) that activates the PDE; the resulting drop in cGMP levels leads to closure of the transduction channels (Fig. 4.1).

There is another class of vertebrate photoreceptor in the higher order layers of the retina, called the ipRGCs (intrinsically photosensitive retinal ganglion cells). Phototransduction in these cells appears to involve biochemistry similar to that found in invertebrate photoreceptors, as we explain below. ipRGCs contribute to non-image forming vision, including pupillary responses (Xue et al. 2011) and photoperiodic processes (Provencio et al. 2002). Although the underlying mechanisms of phototransduction in ipRGCs have not been elucidated, there is a growing body of evidence so far consistent with the idea that light responses in ipRGCs are mediated by pathways including melanopsin as photopigment activating a  $G_q$  type G-protein, leading ultimately to the opening of TRP channels.

Remarkably, melanopsin initiated phototransduction is found in photoreceptors from the earliest chordate, amphioxus, and evidence suggests the involvement of a PLC-dependent signaling cascade, as in *Drosophila* (see below) (Ferrer et al. 2012; Pulido et al. 2012).



**Fig. 4.1** Light transduction in vertebrate and *Drosophila* photoreceptors. *Left.* In vertebrates (*rods and cones*), when a photon hits rhodopsin in the disc membrane it undergoes a photoisomerization and couples to a heterotrimeric G-protein (“transducin”), whose  $G_\alpha$  subunit is released and binds to phosphodiesterase (PDE), activating this enzyme. PDE hydrolyses cGMP, the ligand that opens a cationic non-selective cyclic-nucleotide-gated channel (CNG) in the plasma membrane, as a consequence of which the channel closes, generating a hyperpolarizing receptor potential. *Right.* In the microvilli of *Drosophila* photoreceptors, photoactivated rhodopsin couples to a G-protein which activates the enzyme phospholipase C. This enzyme cleaves the membrane phospholipid phosphatidylinositol bisphosphate (PIP<sub>2</sub>), generating inositol trisphosphate IP<sub>3</sub>, diacylglycerol (DAG) and a proton. DAG somehow opens the TRP and TRPL channels, giving rise to a depolarizing receptor potential. Protons may also contribute to channel opening

### 4.3 *Drosophila* Photoreceptors

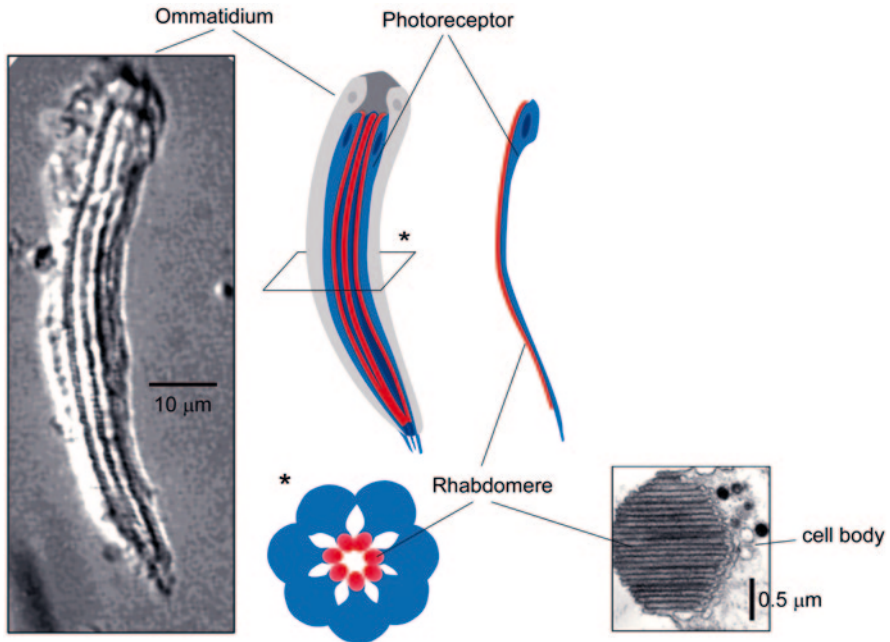
Invertebrate photoreceptors are very different from rods and cones in many ways, biochemically, physiologically and morphologically. There is also great variation among invertebrate photoreceptors, but their photosensitive organelles generally consist of microvilli instead of discs and, with few exceptions, most invertebrate photoreceptors, such as crustaceans, molluscs, arachnids and insects, respond to light with depolarizing receptor potentials resulting from opening of transduction channels, but involving a diversity of signaling cascades.

*Drosophila melanogaster* photoreceptors have been the most extensively studied among invertebrates because they are most experimentally tractable for biochemical, genetic, molecular, physiological and morphological approaches. A diversity of genetic tools and mutants defective in phototransduction have been developed providing an incomparable advantage over the use of other animal species. These tools, in combination with the other techniques, have been effectively utilized for unraveling phototransduction to a remarkable degree.

Studies of *Drosophila* vision led to the discovery of the TRP class of channels, (Montell and Rubin 1989; Cosens and Manning 1969; Hardie and Minke 1992) which has proven to be a central and widespread theme in neural signaling in biology, with particular relevance for many modalities of sensory transduction. Here we review the properties of the two *Drosophila* light-dependent channels, the transient receptor potential (TRP) and the transient receptor potential-like (TRPL), in the context of phototransduction.

The fruit fly compound eye contains some 800 units known as ommatidia, each possessing eight photoreceptors, or retinula cells. Each photoreceptor has a slender cell body ( $40\ \mu\text{m} \times 2\ \mu\text{m}$ ), with a clearly distinguishable longitudinal narrow ribbon-like organelle called the rhabdomere, which consists of roughly 30,000 highly packed photosensitive microvilli. The outer six cells of each ommatidium are positioned circularly beside one another, with two cells positioned centrally; all eight rhabdomeres face each other, oriented towards the inner core matrix of the ommatidium (Fig. 4.2). The axons of the outer six photoreceptors project to the lamina while the axons of the other two central photoreceptors project to the medulla, where they synapse.

Within the microvilli, the phototransduction “machinery” is tightly organized in scaffolded macromolecular assemblies, variously called transducisomes or signalplexes (Tsunoda et al. 1997). The scaffolding protein, INAD, encoded by the *inaD* gene, binds key transduction proteins: TRP, PLC, the regulatory enzyme protein kinase C (PKC) and the myosin NINAC, which anchors the transducing complex to the actin cytoskeleton (Hicks et al. 1996; Wes et al. 1999). INAD scaffolding molecules also bind one another, presumably forming a large supercomplex in the microvillus. Normally, one photon induces only one of these events, termed quantum bump and. Under dim light, it is possible to distinguish the individual quantum bumps, which present obvious differences in size and shape. Using a fly deficient in the regulatory  $\text{Ca}^{2+}$ -binding protein calmodulin, which is necessary for the deactivation of photoactivated rhodopsin, it was observed that a single photon hit triggered multiple repetitive unitary responses (Scott and Zucker 1998). Remarkably, the size and shape of the repetitive events were almost identical, in contrast to those observed in wild type photoreceptors. This result suggests that in the wild type cell every quantum bump originated in a different microvillus, whereas in the mutant the bumps were generated in the only photoactivated organelle where the unitary delivered photon hit. The fact that the single photon response in the mutant consisted of a sequence of individual bumps rather than a continuous depolarization occurred in response to a single photon, is thought to reflect a refractory period following each bump during which the supply of  $\text{PIP}_2$ , exhausted during the response,



**Fig. 4.2** *Drosophila* photoreceptor. Photograph of a dissociated ommatidium (*left*). Schematic representation of an ommatidium depicting the cell bodies of the individual photoreceptors forming part of it and their rhabdomeres (*center*). Individual photoreceptor (*center, right*). Cross section of an ommatidium illustrating the arrangement of the photoreceptors (*center, bottom*); the rhabdomeres of each cell is oriented to the core of the ommatidium. Electron micrograph of the cross section of a rhabdomere, where the highly packed microvilli are appreciated (*right, bottom*) (modified from Hardie 2014)

needed to recover in the microvillus. It is implied that every unitary event would recruit the whole transduction machinery of the respective microvillus, consistent with an all-or-none response. It is concluded that rhabdomeral microvilli are the structural and functional units of phototransduction and contain the proteins responsible for light absorption, signaling cascade, channel opening, and response modulation (Scott and Zuker 1998).

Interestingly, in addition to their rhabdomeral location, TRP and TRPL are also found in photoreceptor axonal terminals in the lamina and the medulla neuropiles of *Drosophila*. Astorga et al. (2012) reported that both channels coexist with the voltage-dependent  $\text{Ca}^{2+}$  channel, called cacophony (Cac), in those cellular regions and they contribute to the transient increase in  $\text{Ca}^{2+}$  required for neurotransmitter release, reinforcing the  $\text{Ca}^{2+}$  influx through Cac. In this synapse, TRP and TRPL work as store-operated channels, but their opening appears to be also mediated by PLC as in the rhabdomere (see below), although opening of these channels is not triggered by rhodopsin as this receptor protein is absent at the synapse. There is additionally  $\text{IP}_3$ -dependent release of  $\text{Ca}^{2+}$  from the endoplasmic reticulum. The contributions from all these  $\text{Ca}^{2+}$  sources allows a massive  $\text{Ca}^{2+}$  elevation that is crucial for the

particularly fast synaptic transmission from the photoreceptors that follows the fast kinetics of phototransduction. Indeed, this synapse is of the ribbon type, characteristically associated to rapid transduction events, also occurring in vestibular hair cells (Lenzi et al. 2002).

#### 4.4 Phototransduction

Light-activated rhodopsin couples to a heterotrimeric GTP-binding protein of the  $G_q$  class. The  $G_{qa}$  catalytic subunit, activated by activated rhodopsin, directly activates inositol-specific phospholipase C (PLC), which is bound in the transducosome to the INAD scaffold. PLC cleaves phosphatidylinositol biphosphate ( $PIP_2$ ), a low abundance plasma membrane phospholipid of importance in signaling, giving three products, inositol trisphosphate ( $IP_3$ ), diacylglycerol (DAG), and a proton (Fig. 4.1). The critical requirement for PLC in phototransduction was revealed by the discovery that a null mutation in the gene coding for this enzyme completely abolished the light response (Bloomquist et al. 1988).

It was initially believed that  $IP_3$  released  $Ca^{2+}$  from an intracellular reservoir, possibly the subrhabdomeral cisternae, close to the microvilli somehow caused the opening of the transduction channels in the microvilli. However, two lines of evidence rule out  $IP_3$  as a required messenger in phototransduction. First, elevating  $IP_3$  in the photoreceptors artificially in the dark by photoreleasing caged  $IP_3$  did not affect subsequent responses to light; and second, mutants lacking  $IP_3$  receptors nonetheless respond normally to light. This leaves the substrate  $PIP_2$  of PLC and the two other products, DAG and  $H^+$ , as possible activators.

The first demonstration that lipids could activate the light-dependent channels was provided by Hardie and collaborators (Chyb et al. 1999). They showed that polyunsaturated fatty acids (PUFAs) added extracellularly to *Drosophila* photoreceptors in darkness generated an inward current with the same current-voltage relation as that activated by light, suggesting that light and PUFAs activated the TRP/TRPL-dependent current. PUFAs are products of the enzymatic activity of DAG lipase, suggesting this enzyme as a key component of the transduction mechanism and therefore present in the microvilli. Nevertheless, immunohistochemical studies that have detected DAG lipase in photoreceptor cell bodies failed to find it in rhabdomeres (Leung et al. 2008). This result, however, must be taken with caution because of the limitation in the resolution of the technique. Several PUFAs were tested by exogenous application to photoreceptors and found to differ in their potency to activate the channels (Chyb et al. 1999). The profile of lipids present in the rhabdomere was unknown at the time, but the presence of ten DAG species with different fatty acid chains, 11 of which are polyunsaturated, namely FUFAs, was recently reported (Muñoz et al. 2013). It was also found that PUFAs not existing endogenously could open TRP channels. Arachidonic acid, which is absent in *Drosophila* rhabdomeres (Muñoz et al. 2013), was able to open the TRP channels. PUFAs exhibit limited solubility in aqueous solutions and partition into the lipid



bilayer, where presumably they induce channel opening. This action could in principle be exerted by directly interacting with the channel proteins or another protein associated with them, or indirectly through transient modifications the structure of the bilayer; however, the mechanism of TRP channel activation is unknown. Subsequent works addressing the role of PUFA in opening the channels have produced conflicting results (Hardie 2014; Muñoz et al. 2013; Delgado et al. 2014).

The other two lipid products of PLC catalytic activity,  $\text{PIP}_2$  and DAG, were extensively investigated by several laboratories.  $\text{PIP}_2$  was first considered on the basis of study on TRPL ectopically expressed in Sf9 insect cells (Estacion et al. 2001). Adding PLC to inside-out excised membrane patches stimulated TRPL activity, while applying  $\text{PIP}_2$  had the opposite effect. These conditions resembled the light and dark situations in the rhabdomere; so, presumably  $\text{PIP}_2$  levels are low and channels are open in light, and  $\text{PIP}_2$  levels are high and channels are closed in dark. However, no direct evidence regarding the actual  $\text{PIP}_2$  levels in the Sf9 cells or of the presumed PLC enzymatic activity in those experiments was reported. Subsequent work on  $\text{PIP}_2$  from different laboratories did not give a consistent picture, as they ranged from inhibitory to excitatory action of the lipid, including absence of an effect on TRPL (Lev et al. 2012). A recent report suggested that  $\text{PIP}_2$  might lead to channel opening in a combined action with a drop of pH that accompanies its hydrolysis by PLC, although the observations done in photoreceptors and excised patches from S2 cells expressing TRPL in this paper are contradictory (Huang et al. 2010).

A role for DAG in TRP and TRPL channel opening has also been explored, in cells heterologously expressing TRPL and in inside-out patches from *Drosophila* rhabdomeres and from expression systems. The results, however, varied among the different experimental preparations (Lev et al. 2012). It is noteworthy that the barrier for DAG to flip from the extracellular to the intracellular membrane leaflet, where is normally located, is much larger than for PUFAs, therefore it is not surprising to fail to see a DAG effect on the channels when applied extracellularly, as opposed to PUFAs. Delgado and Bacigalupo (2009) first showed that DAG applied to inside-out excised rhabdomeral membrane patches of the native membrane was a potent activator of TRP and TRPL. PUFAs could activate the channels as well. A later, more detailed investigation also conducted in excised rhabdomeric patches provided strong support for DAG as the physiological activator of the light-dependent channels (Delgado et al. 2014). With this preparation, the biochemical steps in phototransduction could be reconstituted and manipulated. This study focused on TRP, which is responsible for nearly 95% of the transduction current, whereas TRPL only contributes with 5% (Reuss et al. 1997). When rhabdomeral patches were excised in the dark, TRP remained closed. However, when patches were excised under illumination, TRP was found to be constitutively active, behaving as if the transduction biochemistry in the patch was suspended and the molecular constitution of the patch remained locked in the condition existing at excision. Importantly in this preparation, ATP closed the channels and subsequent application of DAG reopened them; subsequent reintroduction of ATP once more closed the channel. This observation suggested that DAG kinase (DGK), an important component

of phototransduction, was retained in the isolated microvilli membranes and that manipulating its enzymatic activity with ATP mimicked the physiological situation, as this enzyme phosphorylates DAG, generating phosphatidic acid. Although DGK was not detected in rhabdomeres by immunostaining (Masai et al. 1997), the physiological evidence suggests that it is present and functions in DAG turnover (Delgado et al. 2014). This ATP effect was suppressed when DGK was inhibited, either pharmacologically or in the mutant deficient in DGK (*rdgA*). These results strongly implicate DAG as endogenous TRP activator. Evidence militated against the notion of PUFAs as channel activators. If PUFAs were to function as channel activators in phototransduction, functional DAG lipase, which generates PUFA from DAG, would be necessary. If this were the case, one would predict that DGL inhibition would affect TRP opening and that PUFA levels would be light dependent. However, pharmacological inhibition of DGL did not affect the opening of TRP by DAG, showing that conversion of DAG into PUFA was not a necessary step (Delgado et al. 2014). Furthermore, ultra high resolution measurements with liquid-chromatography/mass-spectrometry of DAG and PUFAs levels in a membrane fraction enriched in rhabdomere revealed light-dependent increments in DAG, but no change in PUFAs (Delgado et al. 2014). In this work, PIP<sub>2</sub> and protons were also examined as potential channel activators. Application of PIP<sub>2</sub> to constitutively active excised rhabdomeric membranes patches, in which this lipid presumably was at low level as in darkness, did not alter the channel behavior. Insertion of PIP<sub>2</sub> in the membrane was confirmed by a bioassay in patched from rhabdomeres ectopically expressing the PIP<sub>2</sub>-sensitive Kir2.1 potassium channel. In contrast, introduction of protons did open the channels from the cytoplasmic side of the excised patches. Nevertheless, the effect of acidification from pH 7.15 to 6.4 proved to be irreversible, suggesting that the effect was non-physiological. A previous study had shown that a similar pH change reversible enhanced S2-expressed TRPL basal activity in excised patches, although this was not consistent with experiments in photoreceptors, where pH decrements with bright light were minimal (Huang et al. 2010).

Aside from *Drosophila*, very few other invertebrate photoreceptors have been studied. Scallops have two separate retinas, one containing microvillar and the other ciliated photoreceptors. Although the molecular nature of their light-dependent channels remains unknown, the transduction cascade in the former seems to involve PLC, as in *Drosophila* and their receptor potential is also depolarizing (Nasi and Gomez 1992), whereas in the latter there is a cGMP cascade that leads to the activation of K<sup>+</sup>-selective light-dependent channels (Gomez and Nasi 1994). In the ventral eye of the horseshoe crab *Limulus*, which are also rhabdomeric, the light-dependent channels differ substantially from *Drosophila* and scallop in their relatively large and non-selective cation conductance and ion selectivity; additionally, the underlying signaling mechanism appears to be more complex and is under debate, as there is evidence for the involvement of IP<sub>3</sub>-Ca<sup>2+</sup> and cGMP (Garger et al. 2004; Bacigalupo et al. 1991; Payne and Fein 1986). Summarizing, there is strong evidence indicating that DAG serves as endogenous activator of the light-dependent channels, while PIP<sub>2</sub> and H<sup>+</sup> are not involved directly, and PUFAs do not participate.

Interestingly, mammalian homologs of TRP such as TRPC2, 3, 6 and 7 have been shown to activate by DAG independently of protein kinase C, its most common target (Estacion et al. 2006; Hofmann et al. 1999; Lucas et al. 2003; Okada et al. 1999; Trebak et al. 2003).

#### 4.5 Mechanism of Activation of TRP and TRPL

Although it is well established that lipids related to PLC activity are constituents of the phototransduction cascade and that DAG seems to be the most likely activator of the TRP and TRPL channels, the biophysical basis of the gating of these channels has been scarcely investigated. One obvious possibility is that DAG binds to the channels in one or more specific sites within the membrane. Another alternative is that lipids may lead to channel gating by means of altering the membrane fluidity or curvature, thereby modifying the lipid-channel interaction resulting in a mechanical gating of the channel mechanism, as shown in a mechanotransduction channel (Perozo et al. 2002). A recent report supports this possibility. In a set of elegant experiments Hardie's group (Hardie and Franze 2012) observed a change in photoreceptor length associated with the light response. Notably, the time course of the light-activated current and the photomechanical response measured with atomic force microscopy were similar. The mutant lacking PLC (*norpA*) was unresponsive to light and did not present mechanical responses as well, indicating that the enzymatic activity of this enzyme was crucial. The mechanical response took place even in the mutant devoid of both light-dependent channels (*trp;trpl*), suggesting that it did not involve the channel activity. Furthermore, incorporating the mechanosensitive channel gramicidin in this double mutant restored the ability to generate electrical light responses to these cells, mediated by the truncated phototransduction cascade.

The light response involves a rapid and massive inward current mainly relying on the TRP channel. However, the fact that this channel is normally largely blocked by  $\text{Ca}^{2+}$  like other  $\text{Ca}^{2+}$ -selective channels poses a paradox. Remarkably, it was recently shown that this divalent open channel block can be alleviated upon light activating the PLC enzymatic activity (Parnas et al. 2009), suggesting that the light-induced transient changes in the lipid composition of the membrane moiety of the channels would somehow unblock the channels allowing dramatic transient increment in its  $\text{Ca}^{2+}$  permeability, being rapidly inactivated in a  $\text{Ca}^{2+}$ -calmodulin dependent fashion.

#### 4.6 Molecular and Biophysical Characteristics of TRP and TRPL

Four protein subunits form the TRP and TRPL light-dependent channels, each possessing six transmembrane domains. In this aspect they resemble  $\text{K}^+$  rather than  $\text{Ca}^{2+}$  channels, which possess a subunit having four identical six transmembrane

domains. The *Drosophila* channels share 40% homology between one another and also have a significant homology with mammalian TRPC channels, which are members of the same “canonical” TRP subfamily. The subunit composition of the light-dependent channels was recently shown to be strictly homomeric (Katz et al. 2013).

TRP and TRPL differ in their cation-selectivity. TRP is highly  $\text{Ca}^{2+}$  selective ( $\sim 100 \text{ Ca}^{2+} : 1 \text{ Na}^{+}$ ) while TRPL is poorly selective for  $\text{Ca}^{2+}$  ( $4 \text{ Ca}^{2+} : 1 \text{ Na}^{+}$ ). Around 90% of the phototransduction current flows through TRP. The relevance of TRPL is still mysterious. Under physiological conditions noise analysis of light-dependent currents recorded from dissociated photoreceptors estimated conductances of 8 for TRP and 35 pS for TRPL (Reuss et al. 1997), whereas they respectively have 58 and 40 pS under low divalent cations conditions.

## 4.7 Concluding Remarks

TRP channels in visual transduction have been under investigation for quite some time, and many channel properties and roles have been revealed that have been of broad importance. The mechanism underlying TRP gating has been elusive, however. Recent evidence points to DAG as activator of TRP channels in the *Drosophila* retina. This finding will have significance for many signaling systems, beyond phototransduction and it will have even greater significance when the molecular mechanisms by which DAG activates TRP are elucidated.

**Acknowledgements** FONDECYT 1140520 (JB), CONICYT Graduate Fellowship 22110957 (YM).

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# Chapter 5

## TRP Channels in Transduction for Responses to Odorants and Pheromones

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**Abstract** Transient receptor potential cation channel, subfamily C, member 2 (TRPC2) is a TRP channel expressed by sensory neurons in the vomeronasal organ. Clearly the TRPC2 channel is critically important for transduction in the vomeronasal sensory neurons. However, it appears that TRPC2 is not the only channel that mediates chemical transduction in the VNO. In addition the transient receptor potential channel M5 (TRPM5) is expressed in approximately 5% of the olfactory sensory neurons in the main olfactory epithelium where it plays a role in responding to intraspecific semiochemicals (chemicals involved in animal communication of the same species). Interestingly, TRPC2 is expressed in rodents but not in humans, whereas TRPM5 is expressed in the human raising the question for future work whether this TRP channel is involved in human semiochemical transduction.

**Keywords** Pheromones · Transduction · Olfaction · Vomeronasal · Semiochemical · Ordinary odorants

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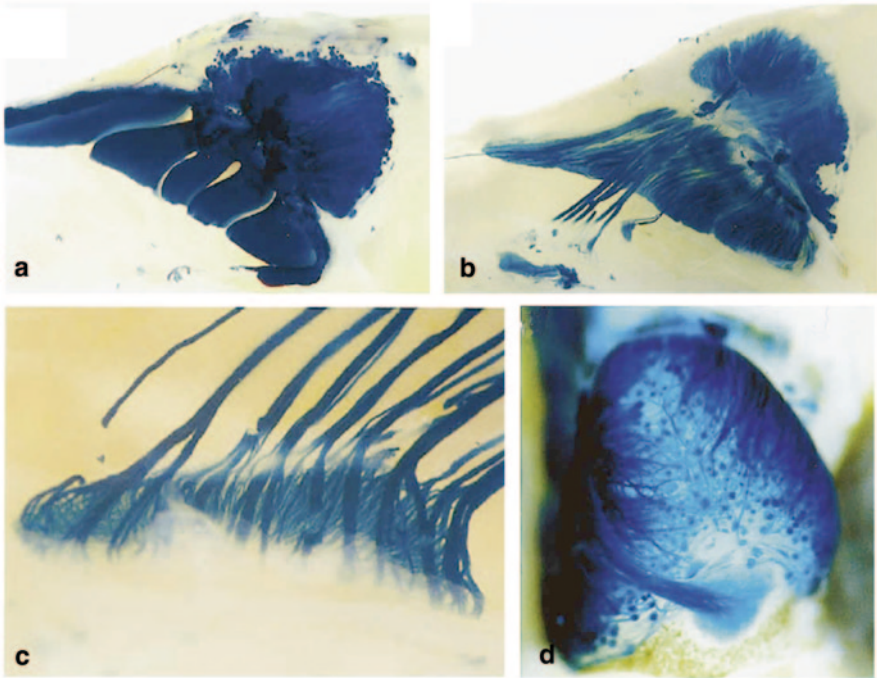
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## 5.1 Introduction

Significant work has determined that the mammalian olfactory and vomeronasal systems possess two chemosensory systems detecting chemicals in the nose and sending information to the main or accessory olfactory bulbs respectively (Fig. 5.1), that express two different TRP channels: TRPC2 and TRPM5 (Liman et al. 1999; Lin et al. 2007). Interestingly, in both systems the TRP channels participate in sensory signal transduction of pheromones and intraspecific semiochemicals. Pheromones were first defined by Karlson and Luscher as ‘substances which are secreted to the outside by an individual and received by a second individual of the same



**Fig. 5.1** Staining of olfactory and vomeronasal sensory neurons blue in the nose and olfactory bulb with X-Gal in OMP-tau-lacZ mice (OMP is olfactory marker protein, expressed in sensory neurons in both VNO and MOE). **a** Whole mount view of the wall of the nasal cavity and the medial aspect of the olfactory bulb. Cell bodies, dendrites and axons are stained intensely such that the epithelium (*left*) and the bulb (*top right*) appear deeply blue. **b** Whole mount view of the nasal septum and the medial aspect of the bulb. The nasal septum, which divides the nasal cavity along the midline, contains three olfactory structures: the main olfactory epithelium (*right*), the vomeronasal organ (*bottom left*), and the septal organ (visible as a small triangle in between the former two structures). The bulb is at the *top right*. Orientation of **(a)** and **(b)**: anterior to the *left*, dorsal at *top*; the *right* half of a mouse head is shown. **c** Whole mount higher power view of the vomeronasal organ. **d** Whole mount view of the dorsal-posterior aspect of the olfactory bulb. Orientation of **(d)**: medial to the *left*, anterior at *top*. Fascicles of blue axons from the main olfactory epithelium terminate in distinct glomeruli in the bulb. Details on the gene-targeted mouse and staining are in Mombaerts et al. (1996)

species, in which they release a specific reaction' (Karlson and Luscher 1959). However, defining mammalian pheromones in this way is problematic, as it is difficult to define what constitutes a 'specific reaction' (Brennan and Keverne 2004). In addition, "semiochemicals" is a broader definition of chemicals involved in animal communication (from the Greek semeion for "sign" that include pheromones) that do involve pheromones. Semiochemicals are different from 'ordinary' odorants.

Importantly, axons from the vomeronasal sensory epithelium whose receptor neurons express TRPC2 target the accessory olfactory bulb that was thought to exclusively send information on pheromones to the vomeronasal medial amygdala (Dulac and Torello 2003; Halpern and Martinez-Marcos 2003; Baum 2012; Baum and Bakker 2013). The medial amygdala then conveys information to certain hypothalamic areas where control of neuroendocrine functions and bisexual social behavior is regulated (Witt and Hummel 2006; Baum and Bakker 2013). However, studies of behavior and glomerular responsiveness indicated that the main olfactory system whose receptor neurons express TRMP5 (Lin et al. 2007) responded not only to ordinary odorants but also to pheromones (Teicher et al. 1980; Dorries et al. 1995; Lin et al. 2008b). Recently it has become clear that a subset of mitral cells in the main olfactory bulb send axons to the vomeronasal amygdala (Pro-Sistiaga et al. 2007; Kang et al. 2009, 2011; Thompson et al. 2012) (see review in (Baum 2012)), implying that a subset of main olfactory system olfactory sensory neurons (OSNs) detect pheromones to convey information to the vomeronasal amygdala.

Here we review recent work on the role of the vomeronasal TRPC2 and main olfactory TRPM5 channels in mediating responses to semiochemicals in the olfactory and vomeronasal systems with a focus on their role on transduction. The reader should refer to other reviews on regular and other aspects of main olfactory transduction (Schild and Restrepo 1998; Munger et al. 2009; Ma 2010) and olfactory and vomeronasal receptors (Axel 2005; Buck 2005; Liberles and Buck 2006; Zhang and Firestein 2009). Finally, this manuscript mentions, but is not focused on other chemosensory cells in the nose and vomeronasal organ that express TRPM5 but are not either OSNs or vomeronasal sensory neurons (VSNs) (Finger et al. 2003; Lin et al. 2008a; Ogura et al. 2011; Tizzano and Finger 2013; Kusumakshi et al. 2015). Importantly there are complementary roles in mediating pheromonal responses for the vomeronasal and main olfactory systems (Brennan and Keverne 2004) and we review below recent work indicating that TRPM5 in the main olfactory neurons responds to semiochemicals.

## 5.2 TRP Channels and the Vomeronasal Organ

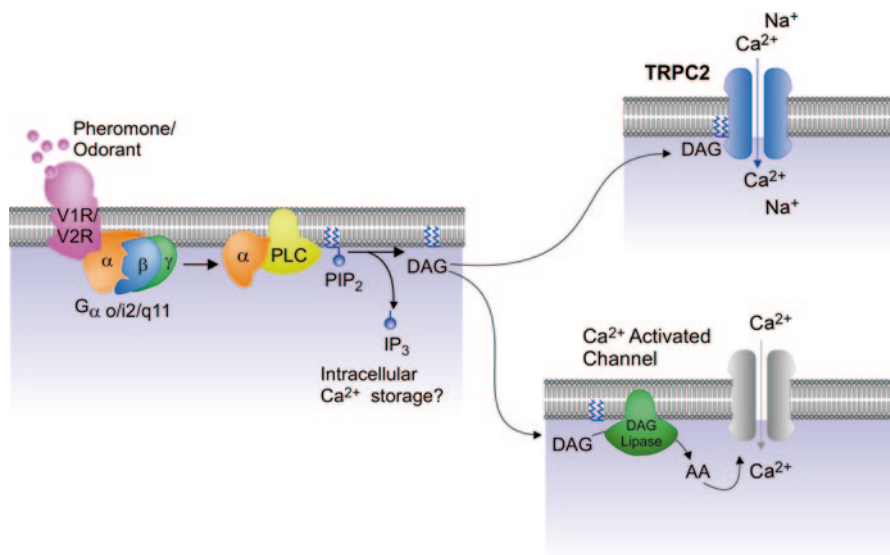
In 1987, Nakamura and Gold showed that  $[cAMP]_i$  directly gates a conductance in olfactory cilia (Nakamura and Gold 1987) and it was subsequently demonstrated that regular odors induce increments in cAMP (Breer et al. 1990) and that cAMP mediates the electrical response (Lowe et al. 1989), implicating that  $[cAMP]_i$  directly gated a conductance that depolarized the olfactory sensory neuron (Nakamura and Gold 1987). Since then substantial work has shown that the activation

of conductance was mediated by a cyclic nucleotide-gated non-selective (CNG) channel (Schild and Restrepo 1998; Munger et al. 2009; Ma 2010). The vomeronasal organ is part of the accessory olfactory system and the VSNs are similar to olfactory sensory neurons in morphology, with the substantial exception that the knobs of the VSNs are crowned with microvilli rather than cilia (expressed by most of the sensory olfactory neurons) and their axons project to the accessory olfactory bulb rather than the main olfactory bulb. However they differ in their transduction pathways compared to sensory neurons from the main olfactory epithelium. Liman and Corey found no evidence for CNG channels in VSNs (Liman and Corey 1996), motivating further research implicating the TRPC2 channel (see below).

### 5.2.1 TRPC2

The TRPC2 channel is highly expressed in the vomeronasal organ although it has been found in lower levels in a few other tissues (i.e. sperm (Vannier et al. 1999; Jungnickel et al. 2001), for review see (Yildirim and Birnbaumer 2007)). TRPC2 was found to be a critical component of the odor transduction cascade in vomeronasal sensory neurons (VSNs) (Liman et al. 1999; Liman and Innan 2003; Brennan and Keverne 2004; Kiselyov et al. 2010). Interestingly, the gene for this channel has been lost in old world monkeys and human genomes but it is robust in most other mammals tested (Liman and Innan 2003; Zhang et al. 2003a; Bacigalupo 2014); moreover, in these animals the vomeronasal organ is vestigial.

The mechanism proposed for activation of TRPC2 in VSNs is similar to that involucrated for the signal cascade in photoreceptors of *Drosophila* that involves a cascade of diacylglycerol that leads to aperture of channels TRP y TRPL (Minke and Parnas 2006; Hardie 2007). In VSNs the canonical signaling pathway starts with a chemical cue binding to a specific G-protein coupled receptor on the microvillar membrane. Once bound, the receptor undergoes a conformational change, activating one of two tightly coupled G protein receptors with  $G_{ai}$  expressed in the VSNs with soma closer to the apical side of the epithelium (V1R) and  $G_{ao}$  expressed in VSNs with cell bodies closer to the basal side of the epithelium (V2R) (Berghard and Buck 1996). The G proteins then activate phospholipase C (PLC) releasing diacylglycerol (DAG) and inositol triphosphate ( $IP_3$ ) (Cinelli et al. 2002; Runnenburger et al. 2002; Iwata et al. 2013). DAG directly gates the closely associated TRPC2 channel, allowing  $Na^+$  and  $Ca^{2+}$  to enter the cell (Lucas et al. 2003; Zufall 2005).  $Ca^{2+}$  influx then activates a  $Ca^{2+}$ -dependent  $Cl^-$  current, leading to further depolarization of the VSN (Yang and Delay 2010; Kim et al. 2011). In a subset of VSNs, DAG is converted to arachidonic acid (AA) by DAG lipase and activates a  $Ca^{2+}$  permeable channel different to the CNG, not yet identified (Spehr et al. 2002; Zhang et al. 2010). Depending on the cell, the species and the location of the smooth ER,  $IP_3$  can enhance the semiochemical-induced  $Ca^{2+}$  transient through release of  $Ca^{2+}$  from intracellular stores (Taniguchi et al. 1995; Inamura et al. 1997; Iwata et al. 2013) (Fig. 5.2).



**Fig. 5.2** Proposed signal transduction of odor responses in VSNs. Pheromone or Odorant (urine) stimulation activate the PLC pathway and elevates DAG, which is converted to AA by a DAG lipase. DAG activates TRPC2, whereas AA activates a Ca<sup>2+</sup>-permeable channels. Both pathways play the excitatory role in the odor responses

The critical need for the TRPC2 channel in VSN signal transduction became clear in 2002 when two seminal papers, by Leybold et al. and Stowers et al. were published (Leybold et al. 2002; Stowers et al. 2002). Stowers et al. (2002) used genetic ablation to create a TRPC2<sup>-/-</sup> mouse (TRPC2 was then named TRP2). Using a sheet of the TRPC2<sup>-/-</sup>VNO neuroepithelium and a multi-electrode array on the microvillar surface, they showed no apparent change in the firing pattern when dilute urine was applied in the knockout compared to baseline. Further, in behavioral studies of male mice, they observed a **decrease** in male/male aggression, using a resident male/intruder male paradigm. Briefly, under this testing procedure, an isolated male mouse established in its home cage has an intruder mouse introduced. Normally, under these conditions, the resident mouse attacks the intruder. When the resident male mouse was a TRPC2<sup>-/-</sup> such aggression did not occur. Rather the TRPC2<sup>-/-</sup> male would often mount the intruder instead. Leybold et al. (2002) deleted the fragment of the TRPC2 channel that contained the transmembrane domains 3–6 and replaced that with a *neo* gene. Their TRPC2<sup>-/-</sup> showed a similar loss on male aggression. In addition, TRPC2<sup>-/-</sup> lactating females showed a decrease in aggression towards an intruder male. They recorded local field potentials of the VNO sensory epithelium and they found the responses to stimuli such as dilute urine were greatly reduced **but not completely eliminated**. Both groups reported that the TRPC2<sup>-/-</sup> males still mated with females and their general behavior seemed normal.

At this point it was clear that TRPC2 was required for many of the VSN responses, but the need for DAG to activate the channel wasn't established until Lucas et al.

(2003) did experiments that showed it was DAG and not  $IP_3$  that activated the channel (Lucas et al. 2003; Zufall et al. 2005). This was important since activation of the original TRP channels discovered was also mediated by DAG (Hardie and Postma 2008; Bacigalupo 2014; Delgado 2014). Further the TRPC2 channels are highly expressed in the dendritic knob/microvillar region of the VSNs (Zufall et al. 2005).

Further research with TRPC2<sup>-/-</sup> female mice has revealed that more than aggressive behavior has been altered. In addition to a reduction in maternal aggression and lactating behavior, these mice show more male type behavior in the presence of a male (Kimchi et al. 2007; Hasen and Gammie 2009). For example, TRPC2<sup>-/-</sup> females will mount males make ultrasound vocalizations and investigation of the anal region of other mice (Kimchi et al. 2007). Interestingly not all responses mediating through the VNO are abolished in TRPC2<sup>-/-</sup> mice (Baum 2012). In pregnant females, the introduction of a strange male results in pregnancy block, called the Bruce effect. If the VNO is surgically removed this effect is abolished (Bellringer et al. 1980). However TRPC2<sup>-/-</sup> females still exhibit the Bruce effect. Research by (Kelliher et al. 2006) suggests that in the basal zone of the VNO VSNs of TRPC2<sup>-/-</sup> mice are still functional and could account for the retention of the Bruce effect. Since only the TRPC2 channel was eliminated it is still possible for DAG to be broken down by DAG lipase to AA and AA to activate a second  $Ca^{2+}$  permeable channel (Spehr et al. 2002; Zhang et al. 2010). Finally, Liman found in VNO neurons a calcium-activated cation channel permeable for  $Na^+$  and  $Ca^{2+}$ , and blocked by ATP and cAMP, that may amplify the primary sensory response (Liman 2003).

Clearly the TRPC2 channel is **critically important** to the functioning of the VSNs, although it appears it is not the only channel that mediates chemical transduction in the VNO (Baum 2012).

## 5.2.2 TRPM5

The TRPM5 channel is a well-known component of taste transduction for Type 2 taste cells. When a tastant binds to its G-protein coupled receptor on the cell membrane it activates PLC leading to an increase in intracellular  $Ca^{2+}$ .  $Ca^{2+}$  in turn activates TRPM5 channels depolarizing the cell (Zhang et al. 2003b; Damak et al. 2006; Liman 2007). In the vomeronasal organ Kaske et al. (2007) reported that the microvillar region of the sensory epithelium appeared to bind antibodies directed at TRPM5 (Kaske et al. 2007). While the authors showed their antibodies were selective for TRPM5 in taste tissue, it wasn't clear if the antibody would bind to TRPC2 but they raised the interesting possibility that the TRPM5 might be involved in VSN transduction. Ogura et al. (2010) explored the role TRPM5 might play in the VNO using a TRPM5-GFP mouse model. They found expression of TRPM5 in the duct of the VNO that they showed to be in a distinct cell subpopulation termed solitary chemosensory cells. They further showed that these cells, when activated, restricted the access of chemicals into the lumen of the VNO (Ogura et al. 2010) Finally, Cre drives expression of tauGFP in TRPM5-IRES-Cre mice in a subset of VNO sensory neurons (Kasumakshi et al. 2015).



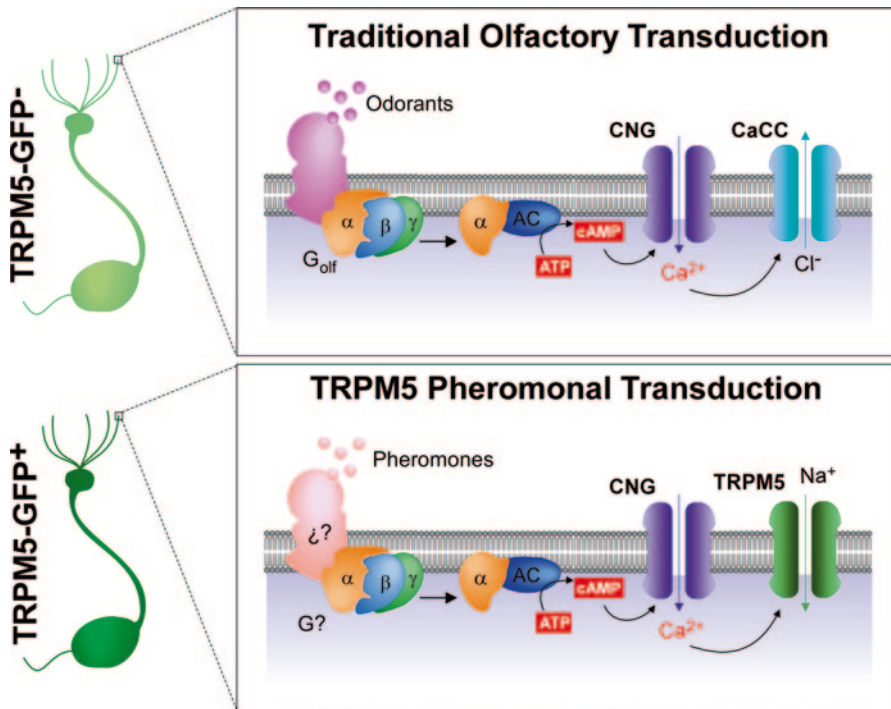
### 5.3 TRPM5 in the Main Olfactory Epithelium

#### 5.3.1 *Ciliary TRPM5 is Responsible for Olfactory Transduction of Responses to Semiochemicals*

Olfactory sensory neurons in the main olfactory epithelium extend cilia (~200 nm wide, tens of  $\mu\text{m}$  long) into the mucus layer overlaying the epithelium (Menco 1997). In the cilia odorants bind to seven transmembrane spanning receptors (Zhang and Firestein 2002; Axel 2005; Buck 2005; Liberles and Buck 2006; Zhang and Firestein 2009) that trigger a G-protein ( $G_{\text{olf}}$ ) (Jones and Reed 1989) to activate adenylyl cyclase III (Bakalyar and Reed 1990) (see Fig. 5.2). Rapidly (within tens of milliseconds (Breer et al. 1990)) the cyclase generates cAMP, which directly opens an unselective cationic cyclic nucleotide-gated channel (CNG) mediating a  $\text{Ca}^{2+}$  influx (Nakamura and Gold 1987; Frings et al. 1995; Baker et al. 1999).  $\text{Ca}^{2+}$  in turn gates a  $\text{Ca}^{2+}$ -activated  $\text{Cl}^-$  channel (Kleene and Gesteland 1991; Lowe and Gold 1993), suggested to be ANO2 (Stephan et al. 2009; Stohr et al. 2009; Rasche et al. 2010) (but see (Gonzalez-Silva et al. 2013)). The electrochemical gradient for  $\text{Cl}^-$  in the cilia favors an efflux of this anion through the  $\text{Ca}^{2+}$ -activated  $\text{Cl}^-$  channel, which significantly potentiates the depolarizing effect of the CNG channel (Kurahashi and Yau 1993; Lowe and Gold 1993) (Fig. 5.3) (but see (Billig et al. 2011)). See reviews in (Schild and Restrepo 1998; Munger et al. 2009; Ma 2012)). The expression of TRPM5 in a subset of OSNs has been suggested by immunohistochemistry, abolished by TRPM5 knockout, and TRPM5-promoter driven expression of GFP (Lin et al. 2007; Oshimoto et al. 2013; Kasumakshi et al. 2015).

Importantly, calcium imaging showed that response of the TRPM5-expressing OSNs to semiochemicals involves discrete increases in  $\text{Ca}^{2+}$  in microdomains (~0.4  $\mu\text{m}$ ) in the cilia, and complementary patch clamp showed that the cilia express  $\text{Ca}^{2+}$ -activated single channels apparently corresponding to TRPM5, as indicated by their sensitivity to TPPO, a specific inhibitor of TRPM5 (Lopez et al. 2014). Furthermore, loose patch experiments performed in this study indicate that in a small number of OSNs (< 5%) semiochemicals elicit activation of OSNs mediated by TRPM5, and not by ANO2 (Fig. 5.4). Finally, the loose patch response of this subset of TRPM5-expressing OSNs to semiochemicals is abolished by knockout of TRPM5 and by the TRPM5 inhibitor TPPO (Lopez et al. 2014).

Incidentally, this explains why the TRPM5 knockout does not affect electroolfactogram to semiochemicals (Lin et al. 2007) because at the concentrations used they would stimulate both the OSNs not expressing TRPM5 and the small number of OSNs that do express TRPM5. Finally, it is interesting that the OSNs that do not express TRPM5 do respond to putative pheromones at high concentration. Likely what these sensory neurons do is to detect putative pheromones and semiochemicals at high concentration as regular odorants explaining why “pheromones” such as 2-heptanaone and 2,5-dimethylpyrazine are detected by mice as a regular odor at high concentration. Indeed, humans can smell these ‘pheromones’.

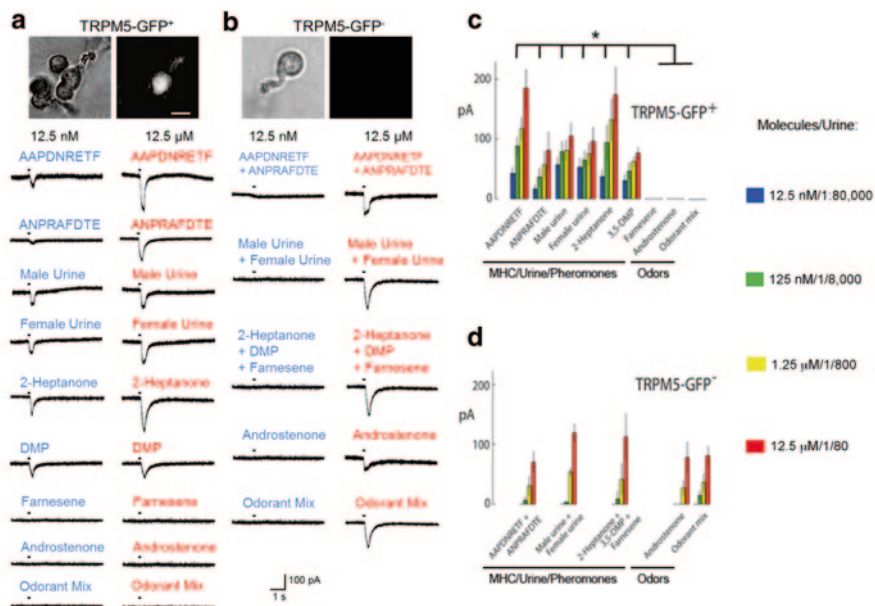


**Fig. 5.3** Diagrams showing: **a** The traditional ciliary olfactory pathway found in OSNs devoid of GFP fluorescence in TRPM5-GFP<sup>-</sup> mice (TRPM5-GFP<sup>-</sup> cells). **b** In this work we show that CNGA2 and TRPM5 are involved in pheromonal transduction in the TRPM5-GFP<sup>+</sup> OSNs. This diagram shows the pheromonal TRPM5 transduction pathway that we propose based on these experiments and the work of Wang and co-workers implicating the involvement of ACIII in pheromonal responses (Wang and Storm 2011). As in odorant transduction both pathways implicate the CNG and a role of  $Ca^{2+}$  entering through this channel activating a  $Ca^{2+}$ -activated  $Cl^{-}$  channel (principally ANO2) and TRPM5 in odorant and pheromone transduction, respectively. Whether  $G_{olf}$  is involved in pheromone transduction in these cells as well, and which proteins are the pheromone receptors remain to be determined. (Reproduced from Lopez et al. 2014)

### 5.3.2 Effect of Knockout of Both CNGA2 and TRPM5

The recent results by Lopez et al. (2014) demonstrate an important route of activating TRPM5 in the OSNs and its role in pheromone signal transduction, which also provides a logic explanation for the co-expression of TRPM5 and CNGA2 in a set of mature OSNs (Lin et al. 2007). However, in the absence of CNGA2 volatile pheromones 2-heptanone and 2, 5-dimethylpyrazine can induce small EOG responses in the MOE and activate glomeruli in the olfactory bulb of CNGA2 knockout mice (Lin et al. 2004). This raises the question whether TRPM5 may be activated independently in the absence of CNGA2. TRPM5 knockout mouse lines were cross-bred with the CNGA2 heterozygous lines to generate offspring that are null to both TRPM5 and CNGA2 (Lin 2014). Clearly, as previously reported (Brunet et al.





**Fig. 5.4** Chemosensory neurons expressing GFP in TRPM5-GFP mice respond to MHC peptides, two pheromones and urine, but not to general odorants and E- $\beta$ -farnesene. (A and B) Photographs of the TRPM5-GFP<sup>+</sup> (A, top) or TRPM5-GFP<sup>-</sup> (B, top) OSNs under phase contrast (left) and fluorescence optics (right) and results of loose-patch current recordings from TRPM5-GFP<sup>+</sup> (A) or TRPM5-GFP<sup>-</sup> (B) OSNs stimulated with 200 ms puffs of pheromones, MHC peptides, urine or odorants. (C and D) Average peak currents ( $\pm$ SEM) of the responses induced by stimuli at different concentrations in TRPM5-GFP<sup>+</sup> (C) or TRPM5-GFP<sup>-</sup> OSNs. (D). The number of OSNs responding to each odor were: TRPM5-GFP<sup>+</sup>: AAPDNRETf (5 of 14), ANPRAFDTE (3 of 30), male C57BL/6 urine (7 of 18), female C57BL/6 urine (7 of 18), 2-heptanone (5 of 18), DMP (5 of 15), E- $\beta$ -farnesene (0 of 15), androstenone (0 of 15) and odorant mixture (citralva, isoamyl acetate, linal, cineol, pine, black pepper) (0 of 20). TRPM5-GFP<sup>-</sup>: mixture of AAPDNRETf and ANPRAFDTE (6 of 17), mixture of male and female C57BL/6 urine (8 of 20), mixture of 2-heptanone, DMP and E- $\beta$ -farnesene (10 of 26), androstenone (5 of 20) and odorant mixture (7 of 17). The asterisk denotes significant differences in responses to the highest concentration of the stimulus determined using rank sum  $p < 0.0025$  with the  $p$ -value corrected by false discovery rate ( $p < 0.03$ ). Odor concentration for g and h: 12.5 nM, 125 nM, 1.25  $\mu$ M and 12.5  $\mu$ M, except for the urine that was diluted 1/80, 1/800, 1/8000 and 1/80,000. Error bars are SEM. These experiments are from Lopez et al. 2014

1996; Baker et al. 1999; Zhao and Reed 2001) CNGA2 knockout negatively impacts the survival of newborn pups that are null to CNGA2, which results in fewer such males at the weaning age (27% of the total male offspring vs. predicted 50% of the males). The CNGA2 knockout pups are known to have difficulty in locating milk source and die at neonatal stage (Brunet et al. 1996). However, the outcome of TRPM5<sup>-/-</sup>/CNGA2<sup>+/-</sup> female and TRPM5<sup>-/-</sup>/CNGA2<sup>+y</sup> male mating is drastically different (Tables 5.1 and 5.2). This raises the question whether the lack of survival of nulls for both TRPM5 and CNGA2 is due to a role of TRPM5 in olfactory signaling processing in the CNGA2 knockout mouse.

**Table 5.1** Offspring (juvenile) genotyping results from CNGA2 mouse line (Mendel prediction in parentheses)

Mating pair genotypes	♂ CNGA2 <sup>+y</sup>	♂ CNGA2 <sup>-y</sup>	♀ CNGA2 <sup>+/+</sup>	♀ CNGA2 <sup>+/-</sup>
♀ CNGA2 <sup>+/+</sup> X ♂ CNGA2 <sup>+y</sup>	58 (42.75)	21 (42.75)	35 (42.75)	57 (42.75)

The genotyping results were obtained from juvenile mice after weaning at 3–4 weeks of age. Pups that died neonatally were not counted. The number of postnatal mice with different genotypes are given for mating of ♂ CNGA2<sup>+y</sup> with ♀ CNGA2<sup>+/-</sup>. The total number of postnatal mice was 171. Bold are different from ♂ CNGA2<sup>+y</sup> or ♀ CNGA2<sup>+/-</sup>

**Table 5.2** Offspring (juvenile) genotyping results from cross mating of TRPM5 and CNGA2 lines (Mendel prediction in parentheses)

Mating pair genotypes	♂ TRPM5 <sup>-/-</sup> , CNGA2 <sup>+y</sup>	♂ TRPM5 <sup>-/-</sup> , CNGA2 <sup>-y</sup>	♀ TRPM5 <sup>-/-</sup> , CNGA2 <sup>+/+</sup>	♀ TRPM5 <sup>-/-</sup> , CNGA2 <sup>+/-</sup>
♀ TRPM5 <sup>-/-</sup> , CNGA2 <sup>+/-</sup> X ♂ TRPM5 <sup>-/-</sup> , CNGA2 <sup>+y</sup>	24 (16.75)	0 (16.75)	25 (16.75)	18 (16.75)

The genotyping results were obtained from juvenile mice after weaning at 3–4 weeks of age. Pups that died neonatally were not counted. The number of postnatal mice with different genotypes are given for mating of ♂ TRPM5<sup>-/-</sup>, CNGA2<sup>-y</sup> with ♀ TRPM5<sup>-/-</sup>, CNGA2<sup>+y</sup>. The total number of postnatal mice was 67. Bold are different from ♂ TRPM5<sup>-/-</sup>, CNGA2<sup>-y</sup>

## 5.4 Future Work on TRP Channels Involved in the Vomeronasal and Main Olfactory Epithelium

The findings have clearly shown an involvement for TRP channels in the VNO. However, there are future studies necessary. An important question is whether there are other TRP channels in the VNO (and the MOE). In a subset of VSNs, DAG lipase converts DAG to AA and activates a calcium permeable channel (Zhang et al. 2010). Is this channel another type of TRP channel? If so, which one? In addition as indicated above, there is expression of TRPM5 in the VNO. Is it just in the solitary chemosensory cells or is there a subset of VSNs that activate this channel rather than Ca<sup>2+</sup>-dependent Cl<sup>-</sup> channels? What is the underlying TRPM5 transduction mechanism in the VNO? What happens if TRPM5 is genetically removed from specific cells and in these knockouts which behavioral activities are affected? Are they activities for “vomeronasal” behaviors?

Interestingly TRPM5 activity is well known to be steeply dependent on temperatures between 15 and 35 °C, and heat activation of this channel is thought to underlie thermal sensitivity of sweet taste (Talavera et al. 2005) (See Chap. 6 by Talavera in this book). This raises the question whether the temperature of the nose decreases substantially affecting the detection of semiochemicals.

There has been considerable controversy on whether higher primates, including human, communicate using ‘pheromones’ and related semiochemicals (Wysocki and Preti 2004; Baum 2012; Baum and Bakker 2013). Nevertheless it is suspected that they respond to semiochemicals, and yet the vomeronasal system appears to be non-functional in humans (Witt and Hummel 2006; Trotier 2011). In addition, the transient receptor TRPC2 expressed in the vomeronasal sensory neurons in rodents is not expressed in primates (Liman and Innan 2003). Yet, there are data indicating that it is possible that OSNs respond to semiochemicals in humans. Thus, for example, putative human male axillary pheromone enhances females’ rating of male sexual attractiveness as well as their sexual arousal (Wyart et al. 2007; Saxton et al. 2008). In addition, human tears contain a chemosignal suspected to be pheromonal (Gelstein et al. 2011). Finally, it is suspected, but not entirely proven, that androstenedione (AND), a putative pheromone that is detected by a particular chemosensory receptor (Keller et al. 2007), may facilitate the attractiveness of potential male sexual partners to women (reviewed in (Baum 2012; Baum and Bakker 2013)). Importantly, the detection of AND is differential for different sexes at certain but not other concentrations (Burke et al. 2012; Baum and Bakker 2013).

## 5.5 Concluding Remarks

Future work is necessary to determine whether human OSNs respond to semiochemicals and, if these cells do respond to semiochemicals, whether TRPM5 is involved in transduction. TRPM5 is expressed in solitary chemoreceptor cells in humans (Barham et al. 2013), but it has not been studied whether this channel is expressed in human OSNs. If it is demonstrated that humans are sensitive to semiochemicals, it will become important to determine whether social and sexual behaviors are affected by semiochemicals.

**Acknowledgements** FONDECYT 1140520 (JB), DC006828 (WL), DC009269 (WL), DC04657 (DR) and DC006070 (DR).

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# Chapter 6

## TRP Channels as Targets for Modulation of Taste Transduction

Karel Talavera

**Abstract** The sense of taste endows organisms with the ability to distinguish nutritious from potentially harmful food components and to orchestrate adaptive attractive or aversive behaviors. Several members of the Transient Receptor Potential (TRP) family of cation channels have been implicated in the perception of canonical and non-canonical taste modalities, TRPM5 for sweet, bitter, umami and fat, TRPP3/PKD1L3 for sour and TRPV1 for salt and metallic tastes. However, TRPM5 is the only one for which there is consensus on its contribution to taste transduction. Here I review recent findings on the role of this channel in taste perception at the peripheral level. Emphasis is made on reported mechanisms of TRPM5 channel modulation that may have an impact on gustatory transduction. Understanding these mechanisms allows learning about basic taste signaling processes and their modulation (e.g. by temperature and taste-taste interactions), and is essential for the design of optimal therapeutic strategies targeting the gustatory system.

**Keywords** Taste transduction · Sweet · Bitter · Umami · Salt · Sour · TRPM5 · TRPV1 · TRPP3/PKD1L3

### 6.1 Introduction

Animal adaptation requires constant monitoring of the chemical environment. In mammals, chemosensation is based on three main mechanisms, olfaction, taste and chemesthesis. The first two allow the detection and identification of chemical compounds that are airborne or dissolved in the saliva, respectively, whereas the third

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© Springer International Publishing Switzerland 2015  
R. Madrid, J. Bacigalupo (eds.), *TRP Channels in Sensory Transduction*,  
DOI 10.1007/978-3-319-18705-1\_6

127

is based on detection at the level of the skin, mucosa and viscera. The initial step in these processes is the interaction of compounds with specialized receptors expressed in sensory cells. This event is then transduced into electric signals, which are carried by nerve fibers to high centers of the nervous system. In the case of taste, transduction takes place in taste buds located in the tongue and palate. These structures contain four types of taste cells (I–IV) and the endings of gustatory nerves (chorda tympani and glossopharyngeal). Type I cells seem to provide mechanical support to the rest of the structure (Pumplin et al. 1997), play buffering functions that maintain low levels of extracellular ATP (Bartel et al. 2006; Vandenbeuch et al. 2013) and  $K^+$  (Dvoryanchikov et al. 2009) and are thought to contribute to salt taste transduction (Vandenbeuch et al. 2008). Type II cells express G protein-coupled receptors (GPCRs) for molecules that we perceive as sweet, bitter or umami (glutamate). Expression of these receptors is more or less exclusive, in the sense that GPCRs for bitter compounds (T2Rs) are not co-expressed with receptors for sweet and umami (heterodimers of T1R1, T1R2 and T1R3) (Chaudhari and Roper 2010). However, it is important to note that the taste quality of a particular taste receptor cell is not determined by the receptors they express, but rather the nerve fiber which they are connected to (Mueller et al. 2005). Type III cells are stimulated by sour and by salts. Type II and type III cells differ in the way they connect to the taste sensory nerves. Upon stimulation, type II cells release ATP and there is evidence indicating that this may occur through the passage of ATP through pannexin 1 (Huang et al. 2007; Romanov et al. 2007; Huang and Roper 2010) and through CALHM1 channels (Taruno et al. 2013a; Taruno et al. 2013b), but it remains unclear whether these mechanisms are mutually exclusive. Extracellular ATP activates P2X receptors in taste nerves and P2Y receptors in Type III cells (Kinnamon and Finger 2013). The latter are endowed with classical synaptic proteins, as well as with voltage-gated  $Ca^{2+}$  channels. Type III cells, also known as ‘presynaptic’, are thought to communicate with the gustatory nerves via vesicular release of GABA and serotonin (Chaudhari and Roper 2010). Finally, type IV cells, also called basal cells, are considered to be precursors of taste receptor cells. These structural and functional aspects of taste were covered by recent reviews (Feng et al. 2012; Liman et al. 2014; Roper 2014).

Several TRP channels have been implicated in distinct aspects of taste transduction. For instance, TRPV1 has been proposed in multiple studies as mediator of the amiloride-insensitive component of salt taste transduction (Lyll et al. 2004, 2005, 2009; Treasukosol et al. 2007; Katsumata et al. 2008). In addition, TRPP3/PKD2L3 were suggested as sour receptors (Ishimaru et al. 2006; LopezJimenez et al. 2006; Ishimaru and Matsunami 2009). However, negative or inconclusive evidence from knockout mice, and lack of definitive proof for the functional expression of these channels in taste receptor cells, have been raised as doubts for their classification as *bona fide* elements of taste transduction pathways (Roper 2014). In contrast, although many issues remain unclear, TRPM5 has drawn consensus about its role in the transduction of several taste qualities. For these reasons, and given the availability of very recent excellent reviews on this subject (Kinnamon 2012; Nilius and Appendino 2013; Liman 2014; Liman et al. 2014; Roper 2014), this chapter focuses on

the most salient basic properties of TRPM5 and how the mechanisms of modulation of this channel influence gustatory transduction at the level of taste receptor cells.

## 6.2 TRPM5 and its Role in Gustatory Transduction and Modulation

### 6.2.1 *Basic Properties in Heterologous and Native Expression Systems*

Of all TRP channels thought to be involved in taste transduction TRPM5 is the best characterized. TRPM5 was first identified in a differential screening of cDNAs from taste receptor cells and was shown to co-express with molecules known to participate in taste signaling, including  $\alpha$ -gustducin and phospholipase C- $\beta_2$  (Perez et al. 2002, 2003). TRPM5 was originally described as a  $\text{Ca}^{2+}$ -permeable store-operated channel (Perez et al. 2002, 2003), but soon afterwards it was established that it is in fact a non-selective  $\text{Ca}^{2+}$ -impermeable cation channel activated by increase in intracellular  $\text{Ca}^{2+}$  concentration (Hofmann et al. 2003; Liu and Liman 2003; Prawitt et al. 2003).

The effective intracellular  $\text{Ca}^{2+}$  concentration ( $\text{EC}_{50}$ ) for TRPM5 activation has been reported for recombinant channels between 0.7 and 32  $\mu\text{M}$  (Hofmann et al. 2003; Liu and Liman 2003; Ullrich et al. 2005) for mouse and 0.84  $\mu\text{M}$  for human (Prawitt et al. 2003). The reasons for these divergences seem to be related to the degree of desensitization of the channel in different recording conditions. Indeed, Ullrich et al. showed that in the whole-cell configuration the  $\text{EC}_{50}$  was 0.7  $\mu\text{M}$ , whereas in inside-out patches it was 32  $\mu\text{M}$  (Ullrich et al. 2005). This desensitization was found to be  $\text{Ca}^{2+}$ -dependent and partially reverted by phosphatidylinositol-4,5-bisphosphate ( $\text{PIP}_2$ ) (Liu and Liman 2003). The mechanisms underlying the sensitivity of TRPM5 to intracellular  $\text{Ca}^{2+}$  remain poorly understood, although it has been discarded that it is mediated by calmodulin (Hofmann et al. 2003). Intracellular  $\text{Ca}^{2+}$  uncaging experiments revealed that TRPM5 is rapidly activated ( $\tau < 2$  ms) upon a strong increase of intracellular  $\text{Ca}^{2+}$  concentration (Ullrich et al. 2005).

The characterization of the biophysical properties of TRPM5 has been limited. The single-channel conductance has been estimated between 16 and 25 pS (Hofmann et al. 2003; Liu and Liman 2003; Prawitt et al. 2003; Zhang et al. 2007), but detailed properties and modulation of single-channel kinetics have not been reported. It is known though that TRPM5 is a voltage-gated channel, activated by membrane depolarization. Like for other voltage-gated TRP channels, the voltage-dependence of TRPM5 activation is quite shallow, with a slope factor between 53 and 58 mV (Talavera et al. 2008c; Gees et al. 2014). This is suggestive of a low apparent gating valence of  $\sim 0.6$  per channel (Talavera et al. 2005), similar to what has been directly estimated for TRPM8 (Voets et al. 2007). In regard to the kinet-

ics of voltage-dependent gating, it has been shown that current relaxation can be described with a single exponential time course. The resulting time constant has a bell-shaped voltage dependence, which is a classical indicative of faster channel closing and opening at very negative and very positive potentials, respectively (Talavera et al. 2008c; Gees et al. 2014). The relationship between the voltage and  $\text{Ca}^{2+}$  dependencies of gating is not yet known, and no gating models have been put forward. As detailed below, TRPM5 was shown to be also strongly stimulated by heat (Talavera et al. 2005).

Most of the properties of recombinant TRPM5 channels have been confirmed in the native environment of taste receptor cells. These include lack of selectivity between  $\text{Na}^+$  and  $\text{Cs}^+$  as charge carriers, impermeability to NMDG<sup>+</sup> and  $\text{Ca}^{2+}$ , stimulation by increase in intracellular  $\text{Ca}^{2+}$  concentration and membrane depolarization and high sensitivity to heat (Romanov et al. 2007; Zhang et al. 2007).

### **6.2.2 TRPM5 in the Perception of Sweet, Bitter and Umami: Two Knockout Mice Tell Slightly Different Stories**

The role of TRPM5 in taste transduction has been directly tested by evaluating the responses of *Trpm5* knockout mice. The first report, by Zhang et al., utilized mice with a partial deletion of the *Trpm5* gene and retaining the promoter region, which suggest that a large part of the amino terminal of TRPM5 could be still transcribed (Zhang et al. 2003). Such mice were reported not to be different from wild type mice regarding the responses to salt and acid stimuli, but completely insensitive to sweet, bitter and umami stimuli at the level of behavior and gustatory nerve responses (Zhang et al. 2003). Thus, it was concluded that TRPM5 is absolutely required for the transduction of these taste modalities. However, other studies that used knockout mice fully lacking the TRPM5 protein consistently yielded different results. In several separate experimental series it was found that gustatory nerve responses to sweet, bitter and umami stimuli were significantly reduced, but not completely abolished in knockout mice (Talavera et al. 2005, 2008c; Damak et al. 2006). Damak et al. (2006) found that *Trpm5* knockout mice were sensitive to sweet, bitter and umami tastes in a long access two-bottle preference test but not in a brief-access licking test. It was argued that this difference could result from a lower sensitivity of the brief-access test and/or from the influence of post-ingestive taste-independent effects (Damak et al. 2006). Interestingly, it was later found that the sweet-blind *Trpm5* knockout mice from Zhang et al. displayed preference for sucrose in a long-term test due to post-ingestive reward (de Araujo et al. 2008). This preference was associated with calorie intake because the non-energetic sweetener sucralose failed to trigger preference over water in the same test.

The reasons underlying the differences in the results from studies with the two *Trpm5* knockout strains remain unknown. It has been suggested that the lack of response of knockout mice in the study of Zhang et al. could be due to a putative dominant-negative action of the TRPM5 N-terminal protein fragment that is

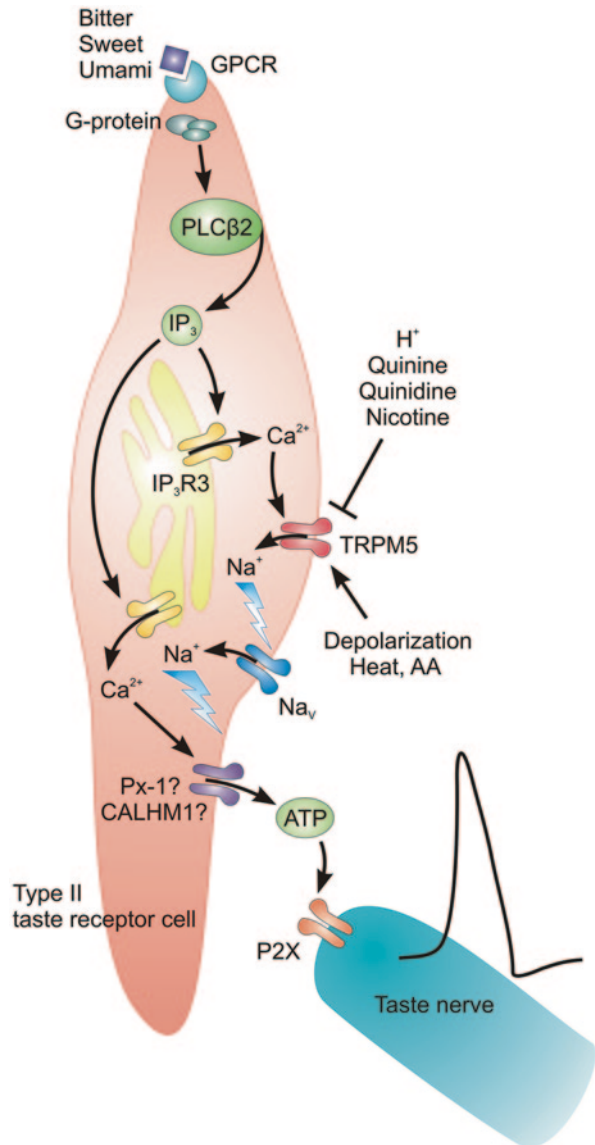
presumably produced in this mice on alternative transduction pathways (Damak et al. 2006). Additionally, differences in genetic background of these mice were also put forward (Damak et al. 2006). However, none of these hypotheses has been tested. Nevertheless, the observations of Damak et al. (2006) and several other studies (Devantier et al. 2008; Eddy et al. 2012 and see below) tilt the balance in favor of a model of transduction of sweet, bitter and umami featuring TRPM5-dependent as well as TRPM5-independent mechanisms. Not surprisingly, the relative importance of these mechanisms depends on the tastant, experimental paradigm and gustatory field being evaluated (Damak et al. 2006).

These and other studies served to establish a qualitative model for the role of TRPM5 in transduction of sweet, bitter and umami stimuli (Fig. 6.1). Activation of G-protein-coupled taste receptors by these tastants stimulates PLC $\beta$ 2 with the consequent production of IP $_3$  and Ca $^{2+}$  release from the endoplasmic reticulum through type 3 IP $_3$  receptors (IP $_3$ R3) (Hisatsune et al. 2007). The increase in intracellular Ca $^{2+}$  concentration activates TRPM5, leading to cation influx and depolarization of the taste receptor cell (Hofmann et al. 2003; Liu and Liman 2003; Prawitt et al. 2003; Zhang et al. 2003). This depolarization activates voltage-gated Na $^+$  channels leading to action potential firing (Gao et al. 2009; Vandenbeuch and Kinnamon 2009). It seems that the combination of increase in intracellular Ca $^{2+}$  concentration and depolarization triggers the release of ATP through pannexin 1 (Huang et al. 2007; Huang and Roper 2010) and/or CALHM1 channels (Taruno et al. 2013a, b).

### 6.2.3 Role of TRPM5 in the Perception of Salts and Fats

Intriguingly, the *Trpm5* knockout mice from Damak et al. (2006) displayed significantly lower responses to NaCl than wild type mice. This again contrasts with the results of Zhang et al. who reported that their *Trpm5* knockout and wild type mice have the same sensitivity to NaCl, but are in agreement with reduced responses to NaCl in  *$\alpha$ -gustducin* (Glendinning et al. 2005) and of *Plc $\beta$ 2* (Dotson et al. 2005) knockout mice. Notably, if we consider TRPM5 to be exclusively expressed in cells transducing sweet, bitter or umami stimuli we should conclude that the TRPM5-dependent responses to NaCl do not elicit salty taste. Instead, they should elicit sweet, bitter and/or umami qualities. Alternatively, TRPM5 may participate in the transduction of saltiness of NaCl in a small proportion of taste receptor cells reported to express this channel but not  *$\alpha$ -gustducin* (Damak et al. 2006) or T1R or T2R (Zhang et al. 2003). This latter possibility is supported by the findings that *Trpm5* knockout mice, but not *Plc $\beta$ 2* knockout mice, display smaller chorda tympani responses to low concentrations (100 mM) of NaCl in the presence of benzamil, a potent blocker of the epithelial Na $^+$  channel ENaC (Ren et al. 2013). On the other hand, the responses of the chorda tympani to salts at high concentrations (500 mM KCl or 500 mM NaCl plus amiloride) were shown to be about 50% weaker in *Trpm5* knockout mice and *Plc $\beta$ 2* knockout mice than in wild type animals (Oka et al. 2013). Interestingly, wild type mice clearly preferred solutions containing

**Fig. 6.1** Cartoon of the TRPM5-dependent signaling pathway for the detection of sweet, bitter and umami stimuli. Type II taste receptor cells express TRPM5 in the basolateral membrane. TRPM5 channels are activated by increase in intracellular  $\text{Ca}^{2+}$  that results from the activation of G-protein-coupled taste receptors sensitive to bitter, sweet and umami compounds. TRPM5 is inhibited by extracellular acidosis, quinine, quinidine and nicotine and stimulated by membrane depolarization, heat and arachidonic acid (AA)



low concentrations of divalent salts such as  $\text{FeSO}_4$  and  $\text{ZnSO}_4$  over water, but this was not observed in *Trpm5* knockout mice nor in *Tlr3* knockout mice (Riera et al. 2009). Furthermore, recent evidence indicates that T1R3 is involved in the detection of calcium and magnesium (Tordoff et al. 2008, 2012).

Finally, there is one study showing that linoleic acid elicits non-selective monovalent cation currents in taste receptor cells that are reduced by application of the TRPM5 inhibitor triphenylphosphine oxide (TPPO) and by genetic ablation of

*Trpm5* (Liu et al. 2011). However, it remains unclear why these maneuvers also reduced the increase in intracellular  $\text{Ca}^{2+}$  concentration triggered by linoleic acid. This is not expected because TRPM5 activation is supposed to be downstream the  $\text{Ca}^{2+}$  signal. Of note, *Trpm5* knockout mice were found to be deficient in the detection of linoleic acid in tests for long-term preference and brief-access licking recordings after conditioned taste aversion. This body of evidence indicates that TRPM5 is involved in the detection of salts and fats, but the precise underlying pathways remain to be fully characterized.

### 6.2.4 Modulators of TRPM5 and their Influence on Taste Transduction

The key role of TRPM5 in transduction mechanisms in the taste receptor cells implies that factors that modulate this channel are expected to affect taste. It is notable, however, that the influences of the  $\text{Ca}^{2+}$ -dependent desensitization and the modulation by  $\text{PIP}_2$  and voltage on taste transduction remain completely unknown.

As other voltage-gated TRP channels (Nilius et al. 2005; Talavera et al. 2008a; b), TRPM5 displays a strong thermal sensitivity, being stimulated by heating. The underlying mechanism entails a shift of the voltage-dependence of channel activation to negative potentials (Talavera et al. 2005), very much like that described for activation of TRPV1 (Voets et al. 2004). Of note, channel stimulation by either depolarization or heating can not be observed at low intracellular  $\text{Ca}^{2+}$  concentration (Talavera et al. 2005). Notably, it was shown that heating strongly enhances chorda tympani responses to sweet compounds in wild type but not in *Trpm5* knockout mice. The strong sensitivity of TRPM5 to heat was therefore proposed to underlie the enhanced sweetness perception at high temperatures. It is also tempting to speculate that heat-induced stimulation of TRPM5 mediates “thermal taste”, which consists in the trigger of taste sensations in the absence of tastants by warming the tongue from 20 to 30 °C (Cruz and Green 2000). This effect would require the stimulation of TRPM5 currents by heating at basal (low) intracellular  $\text{Ca}^{2+}$  concentrations. Alternatively, heating could induce on its own slight increase in the intracellular  $\text{Ca}^{2+}$  concentration and act as initial trigger for  $\text{Ca}^{2+}$ -induced activation of TRPM5. However, none of these possibilities have been addressed experimentally. It has been argued that, besides TRPM5, other elements of the taste transduction pathways could be sensitive to heat and contribute to enhanced taste response (Limman 2006, 2014). However, for any element to be significantly important, it would have to feature a thermal sensitivity similar to that of TRPM5 ( $Q_{10} \sim 10$  between 25 and 35 °C). For any TRP channel lover who recognizes that the “TRP-centric” view of sensory biology has done more harm than good (Meseguer et al. 2011), it would be extremely interesting to identify such heat-sensitive element outside the TRP protein family. At the moment, TRPM5 remains the most plausible candidate (Talavera et al. 2007).

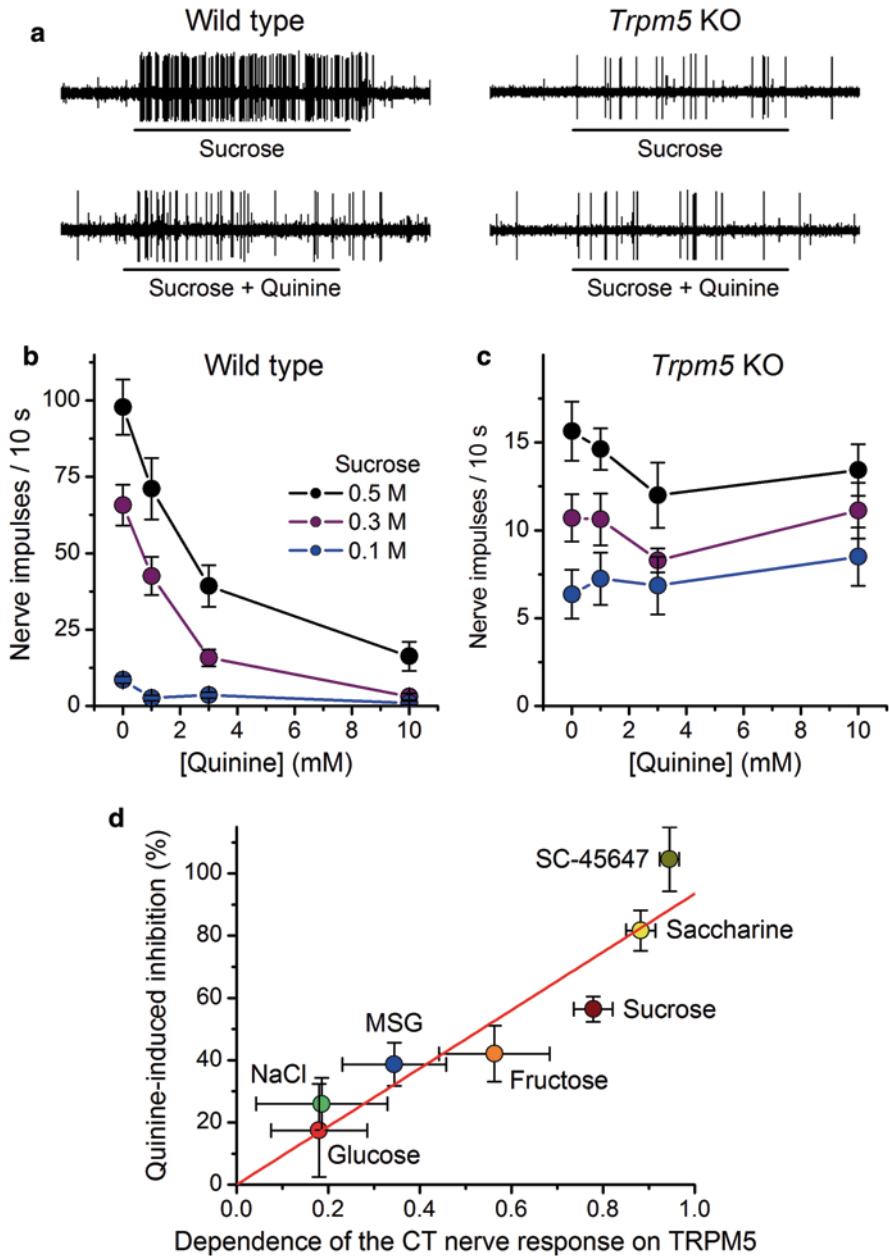


It has been also pointed out that, considering the role of TRPM5 in the transduction of bitter and umami, heat-induced activation of TRPM5 should result in enhanced responses to these stimuli (Liman et al. 2014; Roper 2014). However, this was not observed experimentally (Talavera et al. 2005). To explain this apparent contradiction, it is important to notice that only the responses that strongly rely on TRPM5 activation are expected to be significantly enhanced by heating. In the original report, which used the *Trpm5* knockout strain generated by Damak et al. (2006), only the chorda tympani responses to sweeteners were strongly reduced in the mice lacking TRPM5 (Talavera et al. 2005). As explained below, a similar discussion rose around the inhibitory effect of quinine on TRPM5.

Another potent modulator of TRPM5 is extracellular acidosis. Liu et al. showed that TRPM5 currents were sharply inhibited by drops in extracellular pH between 7 and 5.5 (Liu et al. 2005). This effect was voltage-independent, which is in accordance with the location of residues important for the inhibitory effect at the extracellular face of the channel. Notably, acidosis enhanced the desensitization of TRPM5 currents. Given the restricted localization of TRPM5 at the basolateral membrane of taste receptor cells, the relevance of its inhibition by extracellular protons has been taken with caution (Roper 2014). Also with reserves one should take the reports on stimulation of TRPM5 by arachidonic acid (Oike et al. 2006) and on the inhibition of this channel by extracellular  $Zn^{2+}$  (Uchida and Tominaga 2013). It is currently not clear whether and how these effects are relevant for taste transduction.

The rather extensive literature on interactions between bitter and sweet tastants motivated the study of the possible effects of membrane-permeable bitter compounds on TRPM5 (Talavera et al. 2008c; Gees et al. 2014). It was found that quinine, the bitter additive in tonic waters, and its stereoisomer quinidine inhibit TRPM5 at micromolar concentrations. Furthermore, both compounds inhibited the responses of the chorda tympani to sweet tastants in wild type mice, but did not alter the residual responses of sweet-sensitive gustatory fibers of *Trpm5* knockout mice (Talavera et al. 2008c) (Fig. 6.2). These findings can readily explain the long-known inhibitory effect of quinine on sweet taste perception (Öhrwall 1891) and its partial localization at the gustatory periphery (Formaker and Frank 1996). This phenomenon was considered as a bitter-sweet taste interaction, but it is now clear that the quinine-induced inhibition of the sweet pathway is at least in part dissociated from the detection and processing of the bitter signal because the bitter tastant inhibits directly the sweet transduction pathway.

TRPM5 currents were later found to be inhibited also by nicotine, a compound that has been usually linked to gustatory transduction (Grunberg 1982, 1985; Simons et al. 2006; Lyall et al. 2007; Tomassini et al. 2007). The exact way in which nicotine acts on the perception of these tastes is still not well understood, but very likely multiple different mechanisms are involved. These include nicotine-induced reduction of the size of fungiform papillae and the number of taste cells (Tomassini et al. 2007), inhibition of chorda tympani responses to KCl and NaCl (Lyall et al. 2007), suppression of gustatory responses of neurons in the nucleus of the solitary tract (Simons et al. 2006), inhibition of voltage-dependent sodium channels, sensitization of TRPV1 (Liu et al. 2004) and activation of TRPA1 (Talavera et al. 2009).



**Fig. 6.2** Quinine inhibits the responses of the chorda tympani to sucrose in wild type but not in *Trpm5* knockout mice. **a** Effects of quinine (10 mM) on the responses of single sweet-sensitive fibers from wild type and *Trpm5* knockout mice to sucrose (0.5 M). **b**, **c** Concentration-dependent effects of quinine on the number of nerve impulses elicited by sucrose in single-sweet sensitive fibers from wild type and *Trpm5* knockout mice. **d** Correlation between the inhibitory effects of quinine of the responses of mouse chorda tympani to different tastants with the degree to which these responses depend on TRPM5. (Figure adapted from (Talavera et al. 2008c), with permission)

The inhibition of TRPM5 by nicotine represents yet another peripheral mechanism of inhibition of gustatory perception. Given the relatively high nicotine concentrations required for TRPM5 inhibition ( $IC_{50}$  of 1.3 mM at  $-50$  mV), it is virtually impossible that this effect could be relevant during normal smoking. On the other hand, it could be relevant during tobacco chewing and the use of snuss, in nicotine-replacement therapies, and certainly for the interpretation of experimental studies that used nicotine concentrations of up to 600 mM (Carstens et al. 1998; Boucher et al. 2003; Simons et al. 2006).

Besides shedding light on the mechanisms underlying bitter-sweet taste interactions, the effects of quinine and nicotine on TRPM5 have important implications for our understanding of the mechanisms of bitter taste transduction. The inhibitory effects of these compounds on TRPM5 currents indicate that the perception of their bitter taste cannot be mediated solely by a process requiring TRPM5 activation. Alternative pathways should be recruited to elicit the aversive behavior towards quinine and nicotine at high concentrations. Accordingly, it has been recently shown that nicotine also activates a nAChR-dependent and TRPM5-independent gustatory pathway (Oliveira-Maia et al. 2009). Notably, it is expected that quinine, quinidine and nicotine may also interfere with the perception of umami. However, as for the case of the stimulatory effect of heat on TRPM5 (see above), the effect of TRPM5 inhibitors on the transduction of a certain compound must be related to the extent to which this process depends on TRPM5 activation. This contention was clearly demonstrated for quinine (Talavera et al. 2008c) (Fig. 6.2).

In contrast to quinine, quinidine and nicotine, other bitter compounds known to have an influence in sweet taste perception, including saccharine, caffeine and the related xanthines theophylline and theobromine had no effect on TRPM5 currents, indicating multiple mechanisms of bitter-sweet taste interactions (Gees et al. 2014).

### 6.3 Concluding Remarks

In recent years we have witnessed evident advances in the understanding of the signaling processes underlying taste transduction, but many issues remain unsolved. For instance, it is particularly challenging that despite the large amount of evidence implicating TRPV1 and TRPP channels, no definitive conclusions have been reached. Regarding TRPM5 we are in a more luxurious position, as some tools, including knockout mice, channel modulators and the inhibitor TPPO, are available to study the roles of this channel. We are particularly expectant about the publication of a full report on the discovery and characterization of a group of selective agonists that may help developing new taste modulators and treatments for type II diabetes (Philippaert 2013). However, it is clear that we still know very little about the details of how this channel is gated in basal conditions and how it is modulated. Furthermore, most findings have been obtained in heterologous expression systems and need to be confirmed in taste cells. It is also clear that studying the pharmaco-

logical properties of individual compounds in detail is of outmost importance for the understanding of their specific chemosensory properties.

**Acknowledgments** This work was supported by grants from the Research Council of the KU Leuven (EF/95/010 and PF-TRPLe).

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# Chapter 7

## TRP Channels and Mechanical Transduction

Ana Gomis

**Abstract** Many physiological processes depend on correctly sensing mechanical forces, including hearing, proprioception and touch. Accordingly, much research has focused on the mechanisms and molecules responsible for mechanotransduction. Studies in the fields of genetics, genomics and electrophysiology have converged to further extend our understanding of mechanosensitive events in invertebrates and vertebrates. Indeed, candidate mechanotransduction genes have been identified in mammalian cells, some of which encode the TRP channels expressed in mechanosensitive neurons. In recent years, functional assays have permitted single or multiple ion channel currents flowing through the membrane to be recorded. Such approaches will help determine the biophysical properties of mechanosensitive currents, a crucial step in the quest to identify transduction channels at the molecular level and probe their activity *in vivo*. Here, the proposed mechanisms to mechanodetection are described, along with the different mechanosensory systems used as models to study mechanotransduction. The TRP channels that represent relevant candidates to be involved in sensing mechanical forces will also be reviewed.

**Keywords** TRP channels · Mechanotransduction · Sensory systems · Touch · Pain

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© Springer International Publishing Switzerland 2015  
R. Madrid, J. Bacigalupo (eds.), *TRP Channels in Sensory Transduction*,  
DOI 10.1007/978-3-319-18705-1\_7

141

## 7.1 Principles of Mechanotransduction

In many animals, including humans, some of the most important physiological processes occur through mechanical stimulation. These physiological processes depend on the transformation of mechanical energy into ion currents in order to evoke events such as cellular turgor in bacteria, or the sensation of touch, sound and pain in mammals (Hamill and Martinac 2001; Lewin and Moshourab 2004). To detect and decode mechanical stimuli, organisms possess specialised organs like the touch receptor neurons in *Caenorhabditis elegans*, the chordotonal organ and bristles in *Drosophila melanogaster*, and in mammals, the hair cells in the auditory system, the cutaneous tactile receptors, the circumventricular organs that control systemic osmolarity, the baroreceptors that detect blood pressure, and the proprioceptors that control muscle stretch and the position of the body. In addition, animals also possess free nerve endings of somatosensory and visceral afferents scattered throughout the body (Ernstrom and Chalfie 2002).

As mechanical stimuli are ubiquitous, mechanotransduction may represent one of the oldest sensory processes developed by living organisms. Nevertheless, little is known about the molecular machinery that mediates the sensory detection of force stimuli when compared with our knowledge of other senses.

The mechanisms involved in olfaction, phototransduction and some modalities of taste include molecular mediators that link G protein-coupled receptors to their effector proteins, enzymes or ion channels (Arshavsky et al. 2002; Ronnett and Moon 2002; Hardie and Postma 2008; Kinnamon 2012; Hardie 2014). These mechanisms appear to be evolutionarily conserved and likewise, it could be predicted that there is a general mechanism for mechanical transduction, and a similar sensor that produces different outcomes as a function of its configuration. However, research over the past 10 years indicate that diverse mechanisms and different molecular entities are involved in mechanotransduction, both across species and within single organisms (Sukharev and Corey 2004; Christensen and Corey 2007; Nilius and Honore 2012).

Most mechanosensory cells or mechanoreceptors convert forces into electrical signals, a process that relies on conserved functional principles. The result of this stimulation is the depolarization of the nerve ending and the generation of trains of action potentials (Shepherd 1991; Block 1992; Torre et al. 1995) through the activation of ion channels rather than sensory receptors for odours and most tastes (Garcia-Añoveros and Corey 1997; Sachs and Morris 1998; Hamill and Martinac 2001; Gillespie and Walker 2001). The impulses propagated travel centrally along sensory nerves, eventually reaching cortical structures in order to evoke a sensation.

The idea of mechanically gated ion channels was proposed by Bernhard Katz in 1950 (Katz 1950) on the basis of studies into the muscle spindle. Years later, experiments carried out on hair cells from the bullfrog sacculus (Corey and Hudspeth 1979), bristle mechanoreceptors (Walker et al. 2000), chordotonal hearing receptors from *D. melanogaster* (Albert et al. 2007) and touch receptors in *C. elegans* (O'Hagan et al. 2005) showed that mechanical stimulus produced electrical

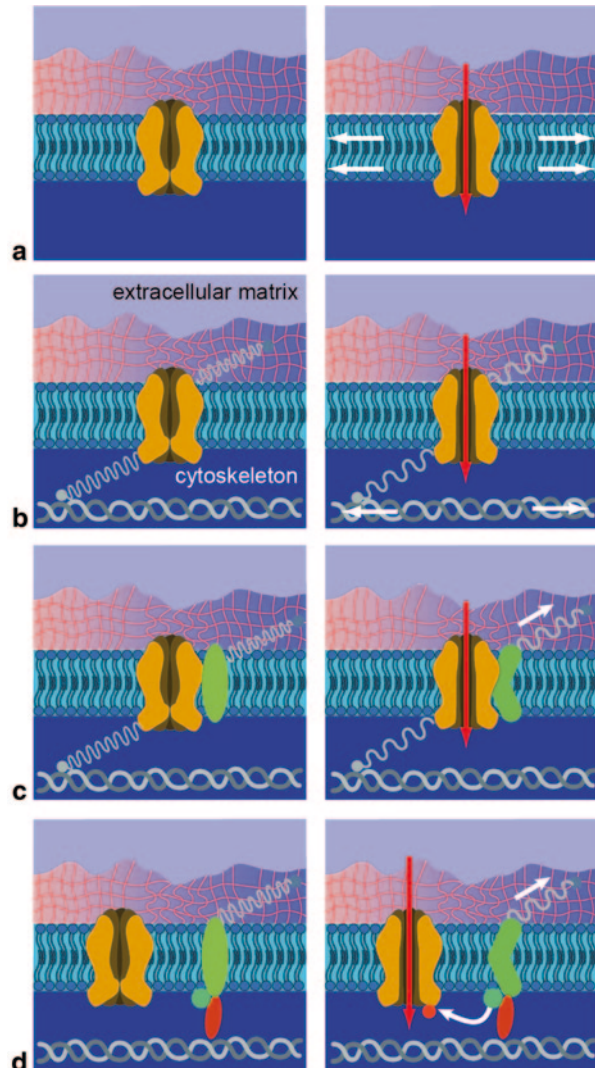
responses in the range of microseconds. This rapid response must result from the direct activation of a transduction channel rather than the activation of second messenger systems. In addition to rapid activation kinetics, the channel must also fulfil other requirements (Ernstrom and Chalfie 2002; Christensen and Corey 2007) : first, the channel must be expressed and located in the receptor cell; second, the candidate protein must be necessary for the electrical signal to be produced in response to mechanical stimulation; and third, the mechanical gating and pharmacological properties of the channel must be similar when it is expressed in different cells.

Studies carried out over the past 15 years have also focused on discovering how these ion channels are activated and although this is still to be fully elucidated, several mechanisms have been proposed to explain how mechanosensitive channels (MS) translate mechanical stimuli into channel opening (Fig. 7.1; Voets et al. 2005; Christensen and Corey 2007; Chalfie 2009; Kung 2005). First, MS channels may be directly gated by the force stimulus. A change in the force may lead to changes in tension of the lipid bilayer, causing conformational changes in the channel and the gating of the pore (Fig. 7.1a). An alternative model for the direct activation of the channel proposes that the channels are 'tied' to cytoskeletal elements and/or the extracellular matrix (ECM), and that the tension among these tethered elements controls the gating of the channels (Fig. 7.1b). However, it is not clear whether these links are directly coupled to the channel domains or if they modulate the forces that surround the channel. This model arises from the biophysical studies in hair cells from the auditory and vestibular systems (LeMasurier and Gillespie 2005) . A third model of activation implicates an accessory protein that will transmit the force to the channel, inducing a conformational change of the channel and indirectly gating its activity (Fig. 7.1c). A fourth possibility is that channels may be activated by intracellular signalling cascades that display mechano- or osmosensitivity (Fig. 7.1d). Mechanical stimulation and osmotic stress evoke numerous changes in protein phosphorylation/dephosphorylation cycles, and it has been shown that some of them play important roles in osmotic and mechanical regulation, acting through protein transport (Pedersen and Nilius 2007).

Multiple mechanisms can converge on a single channel and moreover, the same stimulus may activate different mechanisms. For example, osmotic cell swelling induces tension at the cell membrane that may directly activate channels (mechanism 1) and/or alter the interactions with the cortical cytoskeleton that may in turn affect a tethered channel (mechanism 2). Changes in cell volume may also initiate signalling events that involve the metabolism of membrane components, which may regulate ionic channels by altering the curvature of the membrane (mechanism 4: Pedersen and Nilius 2007) .

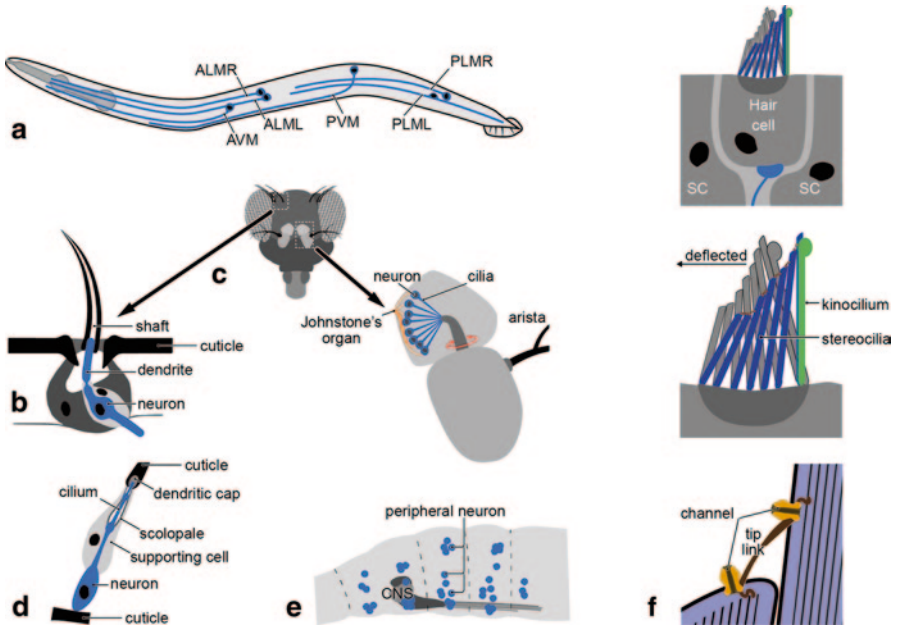
How these mechanosensory mechanisms can be studied will be addressed below.

**Fig. 7.1** Mechanisms of mechanical channel activation. **a** All membrane channels are exposed to lateral forces or tension produce by the modification of the phospholipid bilayer (indicated by the *arrow*), resulting in pore opening. **b** Channel activation through tethered elements, including the cytoskeleton and elements of the extracellular matrix. This interaction may be direct **b** or through proteins associated to the channel **c**. **d** Channel activation through a secondary signal generated by the activation of other proteins sensitive to mechanical forces. (Adapted with permission from Christensen and Corey 2007)



## 7.2 Animal Models in Sensory Mechano-transduction: Mechanosensory Systems

Mechano-transduction is not limited to a subset of specialized cells and tissues but rather, it is implicated in a wide range of cellular functions. The molecular bases of touch sensitivity have been investigated extensively in *C. elegans*. This transparent worm has the particularity of having only 302 neurons, 10% of which are primary mechanosensory neurons, and they have been identified individually through



**Fig. 7.2** Mechanosensory systems. **a** Schematic drawing of *C. elegans* showing the gentle touch neurons. The head of *D. melanogaster* has different mechanosensory organs, bristles (**b**), the Johnston's organ in the antennae (**c**), and chordotonal organs (ch) (**d**). A positive deflection of the bristle stimulates the neurons that innervate it. Sound waves make the arista of the antenna vibrate causing a rotational movement that stretches neurons of Johnston's organ. The characteristic feature of the ch organs is the scolopale cell. Scolopodial receptors are common in tympanal organs and in Johnston's organ, both of which are used in insects to detect external vibration and sound (**d**). Panel (**e**) shows the localization of the multidendritic sensory neurons in *Drosophila* larvae (type II mechanosensory organs). **f** Hair cell and supporting cells (SC) in the sensory epithelium (*upper panel*). On top of the HC the stereocilia and kinocilium in the hair bundle can be observed (*middle panel*). Deflection of the stereocilia would increase the tension in the tip links and open the ion channels (*lower panel*). (Adapted with permission from Christensen and Corey 2007 (**a** and **f**) and from Wilson and Corey 2010 (**b–e**))

laser ablation and genetic studies. Specifically, six of these cells are responsible for light touch and proprioception (Fig. 7.2 a: Chalfie and Thomson 1982). Three cells sense touch in the anterior half of the animal (ALML, ALMR and AVM) and two in the posterior part (PLML and PLMR). Moreover, in addition to touch these cells also appear to regulate other physiological processes (Ernstrom and Chalfie 2002). Other cell seems not to be essential for touch but they mediate movement in touch-mediated circuits (PVM neurons: Chalfie et al. 1985). Over the past three decades, genetic screens have generated a large number of *C. elegans* mutants in which light touch is affected. The relevant proteins involved have been named MEC, from mechanosensory, and they have been numbered from 1 to 18. Two of the most interesting MEC proteins, MEC-4 and MEC-10, form ion channels of the degenerin (DEG) family, which are related to the vertebrate epithelial sodium channels (ENaCs). These two proteins form the pore of the channel, probably together

with MEC-6. This channel associates with other MEC proteins, as well as with the extracellular matrix and cytoskeletal proteins, to form a mechanotransduction complex (for extensive revisions see Garcia-Añoveros and Corey 1997; Ernstrom and Chalfie 2002; Geffeney and Goodman 2012). Genetic studies in worms defective for touch sensation have also revealed the involvement of two related TRPV channels, the OSM-9 and OCR-2 proteins, and of the TRPP2 and TRPML channels (Venkatachalam and Montell 2007).

Studies in the fruit fly *D. melanogaster* have also contributed to our understanding of mechanosensation. Flies display mechanosensation such a gravity sensing, hearing, proprioception, gentle-touch sensation and mechanical nociception. In *Drosophila* there are two types of touch-sensitive cells, the type I cells that have one sensory dendrite (Figs. 7.2b–d) and the type II cells that are multidendritic (Fig. 7.2e; Keil 1997). The most abundant cells are the type I mechanoreceptors, which are associated with accessory structures and that form complex mechanosensory organs. These organs may be associated with external cuticular structures such as bristles (Fig. 7.2b) or cuticular domes (the campaniformsensilla), or they may be attached to the apparently unspecialized cuticle through support cells (the chordotonal organs). The dendrites of these cells terminate at the base of extracellular bristles or domes (external sensory organs) that can be deflected or deformed by touch, airflow or proprioceptive stimulation, or that are surrounded by a supporting cell called a scolopale (Fig. 7.2d). Scolopodial receptors are spindle-shaped cages enclosing an extracellular cavity, into which the ciliar outer segment extends, and they are common in tympanal organs and in Johnston's organ (Fig. 7.2c), both of which are used to detect external vibration and sound (Kernan and Zuker 1995; Eberl 1999).

Type II mechanoreceptors have multiple non-ciliated dendrites, and they lack accessory cells and innervate the insect epidermis (Grueber et al. 2002).

Genetic screening has helped identify genes that are required for the function of the sensory organs in *Drosophila*, whose mutations cause impaired touch responses. One of these genes, named *nompC* (no mechanoreceptor potential C), encodes a channel protein, located in the ciliated mechanosensory organs responsible for bristle touch sensitivity (NOMPC; Walker et al. 2000). This ion channel belongs to the TRPN branch of the TRP channel family (TRPN1) and *NompC* mutants display an approximate reduction in 50% in auditory nerve firing and no amplification of antennal vibration (Eberl et al. 2000), consistent with a hearing deficit. Another gene, *nompA*, was also cloned from Type I cells (Chung et al. 2001), a gene encoding an extracellular linker protein that connects the ciliated dendrites of the auditory mechanosensory neuron to the antennal receiver. Mutation of this gene results in conductive hearing loss. In addition, two more mechanosensory proteins belonging to the TRPV family have been identified, the *nanchung* (*nan*) and *inactive* (*iav*). These proteins are found in the mechanosensory cilia of chordotonal organs, and their mutation results in the loss of auditory responses (Gong et al. 2004; Kim et al. 2003).

Other genetic screens have been performed in *Drosophila* larvae to identify genes responsible for pain responses (pinching and thermal stimuli). In larvae with



deficiencies in the sensation of pain, mutations were identified in a gene (*painless*) that encodes a fly member of the TRPA subfamily channel (Tracey et al. 2003).

More recently, studies on the dendritic arborisation of neurons in *Drosophila* larvae showed that the mechanotransducer channel *piezo* (*Dmpiezo*; see below) is essential for sensing noxious mechanical stimulus *in vivo* (Kim et al. 2012), and that NOMPC is a mechanotransducer channel for gentle touch (Yan et al. 2013). However, a separate study indicates that *Dmpiezo* is not implicated in the mechanotransduction and/or signal amplification for the detection of sound (Zhang et al. 2013).

**Vertebrates** possess a variety of external and internal cells that transduce mechanical stimuli. The external stimuli, touch and sound, are detected by receptors located in the skin and hair cells of the inner ear, respectively. The internal mechanoreceptors in the vascular system, muscle, tendon, organs and joints, and in the urinary bladder detect stretch and pressure stimuli. Among them, the most intensely studied mechanoreceptors are the hair cells and those located in the skin.

**Hair cells** are located in the organ of Corti in the internal ear and they transduce sound and head movements into neuronal signals. The mechanosensory structure of the hair cell consist of a bundle of cilia emerging from the surface of the hair cells (Fig. 7.2f, upper panel), a bundle that contains 30–300 stereocilia and a single kinocilium (Fig. 7.2f middle panel). The stereocilia are connected to each other by a spring protein strand called the tip link, which is thought to be attached to ion channels. The model of mechanotransduction in hair cells proposes that the stimulus produces a deflection of the kinocilia and the associated stereociliary bundle that stretches the tip link. This increase in tension opens ion channels, increasing the inward currents and depolarizing the cells, thereby triggering neurotransmitter release (Fig. 7.2f lower panel) (for extensive reviews see Garcia-Añoveros and Corey 1997; Gillespie and Muller 2009).

Many genes affecting the development and function of hair cells have been identified (<http://www.jax.org/>), some of which were detected due to spontaneous mutation or systematic mutagenesis causing deafness and/or balance defects (Hrabe de Angelis et al. 2000; Nolan et al. 2000; Shearer and Smith 2012). However, identifying the transduction molecules is proving to be more difficult. Evidence has recently supported a direct role for TMC1 and TMC2, members of the transmembrane channel-like family, as components of the transduction complex (Kawashima et al. 2011; Pan et al. 2013; Kim et al. 2013), although it remains uncertain whether TMC1 and 2 are pore-forming subunits and their specific role has yet to be clarified (Morgan and Barr-Gillespie 2013; Holt et al. 2014). Other channels that are candidates to participate in transduction in vertebrate hair-cells are the members of the TRP superfamily, TRPN1, TRPML3, TRPV4 and TRPA1, as will be discussed later on in this chapter.

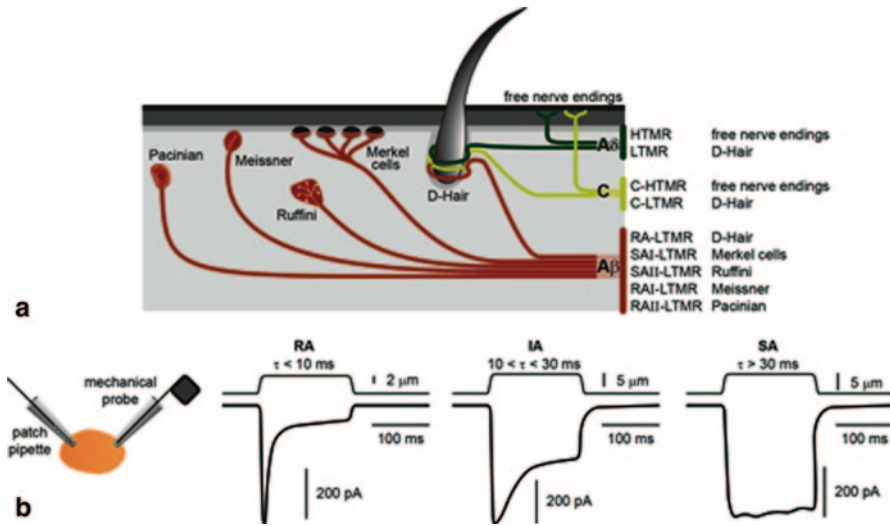
Interestingly, novel mechanical activated currents have been recorded in auditory hair cells lacking tip links. These currents have different properties to those recorded when tip links are functional and the channels responsible is not yet clear. Nevertheless, it has been suggested that normal mechanotransducer channels with an altered distribution or configuration may be involved in producing such currents, as well as ion channels other than mechano-electrical transducer channels, or trans-



duction complex elements other than the TMC1 and TMC2 proteins (Marcotti et al. 2014). Further research should shed light on the identity of these new channels.

**The somatosensory system** in mammals is formed by neurons that innervate the skin, muscle, joints, tendons and internal organs, and that are responsible for the perception and transmission of physical and chemical stimuli originating outside and inside the body. The cell body of these neurons are situated in the dorsal root ganglia (DRG) and trigeminal ganglia (TG), receiving information coming from the body and head, respectively. Transmission of afferent somatosensory information from the periphery to the brain commences with the depolarization of a sensory neuron's terminal, caused by the activation of sensory receptors. These cells include distinct types of mechanoreceptors in function of their anatomy (localization and type of end terminals), their response properties (slow or rapid firing and sensory threshold) or their axonal conduction velocities (myelinated or unmyelinated axons). An important distinction among mechanosensitive neurons is related to the stimulus intensity required to activate their terminals. While some terminals are very sensitive to touch or mechanical displacement, low threshold mechanoreceptors (LTMR), others require stimulation above a high threshold for activation, high threshold mechanonociceptors (HTMRs). Many of the mechanonociceptors are also excited by chemical and thermal stimuli and they are called polymodal nociceptors. Mechanically-evoked nerve impulse discharges are absent in "silent" nociceptor nerve fibres, although they are recorded following local tissue inflammation. The majority of nociceptive fibres are A $\delta$  (myelinated) and C (unmyelinated) type, which terminate with free endings in the tissue they innervate. In terms of the LTMR, the cutaneous receptors are those best known and they can be classified either as A $\delta$  and C fibres whose endings are free nerve terminals, including the D-hair, and as A $\beta$  (myelinated) fibres that terminate in different structures: Pacinian corpuscles, Ruffini endings, Meissner corpuscles (as described in several reviews: Lumpkin and Caterina 2007; Lumpkin et al. 2010; Abraira and Ginty 2013) (Fig. 7.3a). In addition, Merkel's discs have also been shown to function as mechanoreceptors, forming a complex structure with slowly adapting afferents (Maksimovic et al. 2014).

A major issue that is still to be resolved is the identity of the molecules that transduce mechanosensory stimuli, and whether the changes in threshold reflect the activation of different molecular sensors, the differential modulation of the same sensor, or differences in the density or coupling efficiency between the sensor and perireceptor elements. An important reason for the lack of progress regarding these issues is the very small diameter of the mechanosensory terminals and the dispersion of touch receptors in the skin. These factors have prevented the direct recording of mechanogated currents at the site of transduction. In the absence of this type of recordings, different groups have recorded three types of mechanosensory currents from the soma or dendrites of DRG neurons with different kinetics of adaptation: rapidly adapting (RA), intermediately adapting (IA) and slowly adapting (SA: McCarter et al. 1999; Drew et al. 2002; Hu and Lewin 2006; Hao and Delmas 2010) (Fig. 7.3b). Indeed, mechanically activated currents with similar properties have also been recorded in trigeminal neurons (our unpublished results). Some progress



**Fig. 7.3** Cutaneous touch receptors and mechanosensitive currents in mammals. **a** Sensory afferents of neurons innervating skin propagate electrical impulses from the body to the central nervous system. Afferents are classified as  $A\beta$  (red),  $A\delta$  (green) or  $C$  (yellow) based on their cell body sizes, axon diameter, degree of myelination and axonal conduction velocities. Most  $A\beta$  fibres have low mechanical thresholds and  $A\delta$  and  $C$  fibres respond either to noxious stimuli or light touch, each with different physiological profiles and firing patterns (slowly (SA) and rapidly adapting (RA)). The peripheral afferents terminate in different end organs, Merkel cell-neurite complexes, Ruffini endings, Meissner corpuscles, Pacinian corpuscles and hair follicles, and in free nerve ending. **b** Graphical representation of inward mechanosensitive rapidly-adapting, intermediate-adapting and slowly-adapting currents expressed in mice DRG neurons using the mechano-clamp technique (shown in the left panel) at holding potential of  $-80$  mV. Standard mechanical stimuli and time constants of current decay are indicated above the current trace of each panel

has also been made in identifying the mechanotransduction molecules. As such, ASIC 1–3 are expressed in peripheral neurons, although their role in mammalian mechanotransduction is not fully clear (Drew et al. 2004; Roza et al. 2004; Lingueglia 2007). Piezo proteins are pore-forming subunits of a conserved mechanical activated cation family which have recently been considered to be the most promising mechanotransducers candidate (Coste et al. 2010; Coste et al. 2012). In mammals, Piezo1 is necessary for mechanical activated currents in the Neuro2A cell line (Coste et al. 2010) and Piezo2 mediates the mechanical responses in Merkel-cells, evidence of a physiological role in mammalian mechanotransduction (Woo et al. 2014; Maksimovic et al. 2014; Ikeda et al. 2014). Nevertheless, the involvement of TRP channels in mechanotransduction cannot be ruled out, as discussed below.

Mechanotransduction also plays a fundamental role in the **vascular and renal system, and the urinary bladder** among others. These systems are permanently subjected to mechanical forces, such as shear stress and pressure, and impaired handling of these stimuli in these systems can cause severe pathologies. Many studies have focused on the molecular mechanisms of mechanical stretch in visceral

systems and thus, this issue will not be addressed in this chapter (for reviews see Haga et al. 2007; Lehoux et al. 2006; Li and Xu 2007; Orr et al. 2006; Shyu 2009; Hahn and Schwartz 2009; Weinbaum et al. 2010). Mechano-gated ion channels play a key physiological role in the visceral system including Piezo1, that plays a critical role in endothelial cell mechanotransduction (Ranade et al. 2014), and the TRP channels. This family of channels that are expressed in different cell types within the heart, kidney and bladder we will now centre out attention on.

### 7.3 Mechanosensory TRP Channels

The channel proteins that belong to the TRP family appear to mediate many forms of sensory perception, the reason why they are the focal point of the reviews in this book. There is some controversy as to whether mechanosensitive TRP channels are activated directly or if they are specifically involved in mechanical signalling. Moreover, piezo2 was recently shown to be required for Merkel cell mechanotransduction. Still, there is strong genetic evidence that a variety of TRP channels contribute to mechanosensation (Table 7.1). This includes evidence from loss-of-

**Table 7.1** Mechanosensory channels

Molecular identity	Family	Organism	Mechanosensitivity
TRPA1	TRPA	Mammals, worm, fly	Pressure, stretch, osmolarity & touch
TRPC1	TRPC	Mammals/ <i>Xenopus</i>	Touch, stretch, osmolarity & shear stress
TRPC3	TRPC	Mammals	Gentle touch & audition
TRPC5	TRPC	Mammals	Osmolarity & pressure
TRPC6	TRPC	Mammals	Osmolarity, pressure, touch & audition
TRPV1	TRPV	Mammals	Stretch & osmolarity
TRPV2	TRPV	Mammals	Stretch & osmolarity
TRPV4	TRPV	Mammals	Noxious pressure, shear stress, osmolarity & stretch
TRPM3	TRPM	Mammals	Osmolarity
TRPM4	TRPM	Mammals	Osmolarity
TRPM7	TRPM	Mammals	Osmolarity
TRPN1	TRPN	Fly, fish & amphibian	Touch & audition
OSM-9/OCR-2	TRPV	Worm	Osmolarity, audition & touch
TRPP2	TRPP	Mammals, worm, fly	Pressure, osmolarity & shear force
TRPML3	TRPM	Mammals, worm, fly	Audition

function mutations in organisms ranging from invertebrates to zebra fish, and from mice and humans (Christensen and Corey 2007; Eijkelkamp et al. 2013).

### 7.3.1 TRPA1

The ankyrin-repeat channel TRPA1 is strongly conserved in the animal kingdom and it is found in many vertebrates and invertebrates. The *C elegans* TRPA1 is activated by mechanical stimulus, suggesting a role for TRPA1 in mechanosensation (Kindt et al. 2007). Indeed, TRPA1 is required for normal responses to nose-touch, and for a specific foraging behaviour of the worm in the presence of food that consists of high frequency and local exploratory movements of the head. In *Drosophila* larvae that lack *Painless*, the invertebrate orthologue of TRPA1, mechanical nociception is dampened (Tracey et al. 2003). It has been suggested that distinct isoforms of *painless* are responsible for specific sensory modalities that require its function. Moreover, the function of *painless* in noxious mechanosensation is independent of its ankyrin repeats as this phenotype is reverted after the introduction of a short isoform of the *painless* gene in mutant flies (Hwang et al. 2012).

In mammals, TRPA1 is expressed in sensory neurons of the dorsal root, trigeminal and nodose ganglia, and in many organs and tissues, including the brain, inner ear, smooth and skeletal muscle, and pancreas (Nilius et al. 2012). TRPA1 was initially a strong candidate to act as the hair cell transduction channel in vertebrates (Corey et al. 2004) on the basis of three main observations: (i) TRPA1 is expressed in the hair cells of both the cochlea and the vestibular system; (ii) TRPA1 messenger RNA transcripts appear on embryonic day 17, coinciding with the onset of mechanotransduction; and most importantly, (iii) disruption of TRPA1 expression in zebra fish with morpholino oligonucleotides, and in mice with small-interference RNA, strongly inhibits hair cell mechanotransduction. TRPA1 was also found to be expressed in the free nerve endings and in the bundles of stereocilia, leading to the proposal that it fulfils a role in both nociceptive and auditory cell function (Nagata et al. 2005). However, subsequent studies in zebrafish larvae lacking the orthologues of mammalian TRPA1 exhibited normal mechanosensory hair cell function (Prober et al. 2008). In addition, hearing and vestibular function is normal in two different TRPA1<sup>-/-</sup> mouse lines (Kwan et al. 2006; Bautista et al. 2006), although these two lines have a conflicting nociceptive responses. One of these lines exhibited pain hypersensitivity (Kwan et al. 2006), TRPA1<sup>-/-</sup> mice displaying a higher mechanosensory pain threshold than their TRPA1<sup>+/+</sup> littermates as well as a weaker response to supra-threshold stimuli. By contrast, no difference in the mechanical threshold was detected between TRPA1<sup>-/-</sup> and wild type animals in the other line (Bautista et al. 2006).

More evidence of a role for TRPA1 in sensing mechanical stimuli has been obtained through recordings in the skin-nerve preparation from TRPA1<sup>-/-</sup> (Kwan et al. 2009). In the absence of TRPA1 both C and AM fibre nociceptors had a lower firing rate at all force intensities and for noxious force, respectively. TRPA1 was

expressed in many medium to large DRG somata and on large-calibre axons associated to A $\beta$ -fibres, and hence it was concluded that TRPA1 reduced the slow adaptation to sustained force in A $\beta$ -fibres but increased the firing of D-hair and RA A $\beta$ -fibres (Kwan et al. 2009). Moreover, pharmacological inhibition of TRPA1 using the selective antagonist HC-030031 and genetic ablation has shown that cutaneous C fibres require TRPA1 to respond to noxious stimulus evoked by formalin application, as well as to high intensity mechanical forces. However, HC-030031 had no effect on mechanical firing in nociceptive A $\delta$  fibres (Kerstein et al. 2009).

The contribution of TRPA1 to mechanical sensitivity was also assessed by studying the mechanically-gated currents in cultured DRG neurons induced by applying an electrically driven mechanical probe to the cell membrane. TRPA1 deletion in knockout mice significantly reduces the maximum IA currents in small-diameter neurons, producing no effect on the RA and SA currents in small and large neurons. Moreover, the inhibitor HC-030031 significantly decreases IA current amplitude in wild type mice activated by the TRPA1 agonist AITC (Brierley et al. 2011). Similarly, SA currents were absent in IB4 negative neurons from TRPA1<sup>-/-</sup> mice and the amplitude of RA and IA currents in IB4 positive neurons from TRPA1<sup>-/-</sup> mice was reduced by over the 60%. Moreover, HC-030031 was only seen to dampen SA currents wild type mice (Vilceanu and Stucky 2010). Together, it seems that TRPA1 mainly participates in SA and IA currents.

So far, direct evidence of TRPA1 activation by mechanical forces has only been obtained in HEK cells transiently expressing TRPA1, and only in response to a hyperosmotic solution but not to a hypoosmotic solution (Zhang et al. 2008). By contrast, TRPA1 does contribute to the Ca<sup>2+</sup> influx evoked by hypoosmotic stimulation-induced plasma membrane stretching in Merkel cells from hamster buccal mucosa (Soya et al. 2014). TRPA1 also plays a role in the mechanical hyperalgesia and allodynia induced by the gasotransmitter hydrogen sulphide in mice (Okubo et al. 2012) and in the mechanical hyperalgesia caused by a deep-tissue incision in the rat hind paw (Wei et al. 2012). Mechanical hyperalgesia induced by Freund's adjuvant in wild type mice was strongly reduced by blocking TRPA1 with the small molecule, AP18. However, the fact that AP18 does not prevent Freund's adjuvant-induced mechanical hyperalgesia in TRPA1<sup>-/-</sup> mice may suggest that TRPA1 is involved in the maintenance of mechanical hyperalgesia rather than its generation (Petrus et al. 2007).

Several studies point to a role of TRPA1 in visceral mechanotransduction. TRPA1 is expressed in the rodent and human bladder, and activation of TRPA1 by agonists or changes in its expression is associated with bladder dysfunction (Du et al. 2007, 2008; Streng et al. 2008). A role for TRPA1 in noxious colonic distension was also reported (Brierley et al. 2009).

Together, these results suggest that TRPA1 plays a role in mechanosensation although it remains to be determined whether it acts as a mechanical sensor or if it indirectly participates in mechanosensation by amplifying or modulating the signal from the transduction channel.

### 7.3.2 TRPC

Examining the mechanosensitivity of TRP channels in heterologous system has provided conflicting data in some instances. This is certainly the case of **TRPC1** which was first considered to be a mechanosensitive non-selective cation channel (MscCa) given that application of positive and negative pressure resulted in a large increase in current that can be attributed to MscCa in *Xenopus* oocytes and in CHO-K1 cells transfected with human TRPC1 (Maroto et al. 2005). However, a study published 3 years later showed the difficulties of testing mechanosensitivity in heterologous systems due to the background mechanical activity observed in cell lines (Gottlieb et al. 2008).

To date, the only proven role of TRPC1 in mechanosensation was described by studying the recordings of the afferent terminals that innervate the hairy skin of TRPC1<sup>-/-</sup> mice (Garrison et al. 2012). In these animals there was an ~40% reduction in action potential firing of SA-A $\beta$  fibres in response to forces at intensities lower than 4–200 mN yet not in the RA-A $\beta$  fibres. A reduction of 50% in the discharges was observed in the D-hair A $\delta$  fibres, whereas C-fibres and nociceptive AM fibres innervating hairy skin responded normally to noxious mechanical stimulation in TRPC1<sup>-/-</sup> mice, meaning that TRPC1 does not play significant role in cutaneous mechanical nociception. However, in a behavioural light-touch test a significant decrease in mechanical sensitivity was evident in TRPC1 deficient mice compared to wild type mice (Garrison et al. 2012). More recent findings indicate a reduced static firing rate of SA-A $\delta$  fibres and a moderate decrease in the behavioural response to light touch in an epidermal piezo2 knockout mice (Maksimovic et al. 2014). These latter data might suggest that TRPC1 and piezo2 mediate mechanotransduction in Merkel cells.

Within the cardiovascular system, pressure-induced constriction of cerebral arteries in smooth muscle cells is unaffected in TRPC1 knock-out mice, indicating that TRPC1 is not an obligatory component of stretch-activated ion channel complexes in vascular smooth muscle (Dietrich et al. 2007).

**TRPC3** has been implicated in the kinetics of mechanical activated currents, given that the proportion of sensory neurons that display IA currents increased in TRPC3<sup>-/-</sup> mice while those responding with RA currents decreased (Quick et al. 2012). Interestingly, TRPC3 was also attributed a role in the detection of light touch when it forms a complex with TRPC6. Indeed, there were deficits in innocuous mechanosensation in the TRPC3 and TRPC6 double knockout mice but not in the single KO mice. There was also a significant decrease in the proportion of neurons displaying RA currents in these double KO animals, and elimination of TRPC3 and TRPC6 also produced hearing deficits (Quick et al. 2012).

Mechanical activation of **TRPC6** has been addressed, and it was reported that osmotic and pressure-induced membrane stretch directly activates the TRPC6 channel expressed in heterologous systems. The fact that TRPC6 is strongly expressed in vascular smooth muscle cells made it a candidate mechanotransducer in smooth muscle (Spassova et al. 2006). However, later studies demonstrated the activation



of TRPC6 through a Gq/11-coupled protein receptor (Schnitzler et al. 2008; Inoue et al. 2009; Sharif-Naeini et al. 2010). TRPC6 has also been involved in kidney regulation via podocin modulation, a protein involved in the function of the filtration barrier in the mammalian kidney. However, the implication of podocin in mechanosensation is in any case speculative, such that the participation of TRPC6 and other members of the TRPC subfamily is probably indirect (Huber et al. 2006).

**TRPC5** has been proposed to act as a direct sensor of lysophospholipids (Flemming et al. 2006), suggesting that this channel is sensitive to the structure of the lipid bilayer. We characterized the activation of **TRPC5** in response to hypoosmotic stimulation and positive pressure in a heterologous system expressing TRPC5. This activation is blocked by the tarantula peptide GsMTx-4, a specific inhibitor of stretch activated channels, and it is independent of PLC activation (Gomis et al. 2008). Interestingly TRPC5 is also activated by applying force to the cell's surface via a glass pipette controlled by a piezo-electric system, provoking currents with RA kinetics (unpublished data). TRPC5 channels are expressed in trigeminal and DRG neurons, suggesting a possible role in mechanotransduction, although their physiological role in mechanotransduction and osmotic regulation remains to be clarified.

### 7.3.3 *TRPV*

Several TRPV channels have been described to be sensitive to mechanical stimulus or volume/osmotic change. **TRPV1** is required for osmosensory transduction in the organum vasculosum lamina terminalis, a primary osmosensor nucleus in the brain (Ciura and Bourque 2006; Ciura et al. 2011). Although controversial, the TRPV1 channel has been implicated in mechanical hypersensitivity as a consequence of inflammation, heat injury or nerve ligation (Honore et al. 2005; Bolcskei et al. 2005). The reason for the discrepancies observed is not clear but it is possible that TRPV1 may be involved in cutaneous mechanical hyperalgesia through an indirect mechanism rather than as a direct noxious mechanical sensor. The involvement of TRPV1 in mechanical hyperalgesia is more obvious in deeper tissues than in cutaneous pain.

**TRPV2** is expressed in mesenteric and basilar arterial myocytes and it is activated by membrane stretch in TRPV2-CHO cells. Therefore, it would seem that TRPV2 can function as a stretch-activated channel in vascular smooth muscle (Muraki et al. 2003), although studies in intact arteries are still needed to confirm the physiological role of TRPV2. Studies in somatosensory systems indicated that TRPV2 is not essential in mechanical nociception (Park et al. 2011).

**TRPV4** was first reported as an osmotically activated channel over a decade ago, when it was shown that the osmotic stimulus activates the TRPV4 (previously known as VR-OAC or OTRPC4) expressed in HEK and in CHO cells (Liedtke et al. 2000; Strotmann et al. 2000; Wissenbach et al. 2000). However, a subsequent study failed to find significant activation of TRPV4 expressed in CHO cells by hy-



poosmotic stimulation (Suzuki et al. 2003), while the activation of TRPV4 by cell swelling was seen to occur through the PLA2-dependent formation of arachidonic acid and its subsequent metabolism to 5',6'-epoxyeicosatrienoic acid (Vriens et al. 2004). The contribution of TRPV4 *in vivo* was studied by expressing the mammalian TRPV4 in *C elegans* sensory neurons and using TRPV4<sup>-/-</sup> mice. TRPV4 is required for nose touch avoidance in *C elegans* (Liedtke et al. 2003) and its loss markedly reduced the sensitivity of the mouse tail to pressure (Suzuki et al. 2003). Hypotonic solutions activate 54% of C-fibres in rats and the effect of hypoosmotic stimulation is more effective in inflammation states, suggesting that TRPV4 plays an important role in mechanical hyperalgesia (Alessandri-Haber et al. 2003; Alessandri-Haber et al. 2006). Moreover, proteases generated during inflammation cause mechanical hyperalgesia through TRPV4 sensitisation (Grant et al. 2007). Interestingly, interactions between different TRP channels, such as TRPC1, TRPC6 and TRPV4 contribute to the development of mechanical hyperalgesia, although they may fulfil different roles (Alessandri-Haber et al. 2009).

TRPV4 is expressed in hair cells and thus, the role of TRPV4 in the transduction of acoustic stimuli has also been studied. The cochlea of TRPV4<sup>-/-</sup> mice has a normal morphology but these animals develop late-onset hearing loss, which is similar to the autosomal dominant non-syndromic hearing loss in humans, and loss of TRPV4 makes the cochlea vulnerable to acoustic injury (Tabuchi et al. 2005). There is also evidence of a possible role for TRPV4 in skeletal muscle fibres. Recordings of single channel activity showed that hypoosmotic solutions activate mechanosensory channels and the absence of mechanosensory activity in the muscle of TRPV4<sup>-/-</sup> mice (Ho et al. 2012). However, the implication of other TRP channels forming heteromeric proteins with TRPV4 cannot be ruled out.

### 7.3.4 TRPM

Three members of the TRPM subfamily have been implicated in osmosensation: the **TRPM3** channel that is expressed in the odontoblast (Son et al. 2009) and kidney (Grimm et al. 2003); **TRPM4** in blood vessels (Morita et al. 2007) and that has been implicated in cerebral blood flow regulation (Reading and Brayden 2007); and **TRPM7** that is expressed in human epithelial HeLa cells and is thought to be involved in regulation associated with a volume decrease (Numata et al. 2007). Although these data imply an inherent stretch or mechanosensitivity of these channels, further studies are necessary to define the functional role of these channels in mechanotransduction.

### 7.3.5 TRPN1

The *nompC* gene encodes the **TRPN1** channel in *Drosophila melanogaster* and it is involved in touch, hearing and proprioception. Mutations, in this gene result in de-

iciencies in bristle mechanotransduction (Kernan et al. 1994; Walker et al. 2000). However, while TRPN1 is an excellent candidate mechanotransduction channel in flies and fish, *nompC* homologues have not been found in the mammalian genome (Sidi et al. 2003; Shin et al. 2005; Li et al. 2006).

## 7.4 Concluding Remarks

Since their discovery three decades ago, TRP channels have been shown to participate in a number of different sensory processes. Still, our understanding of the functional roles of this ion channel superfamily is limited in comparison to what we know about those of ligand- and voltage-activated channels. Progress included the discovery of the involvement of some of the TRP channels in the detection of mechanical forces, although as discussed previously in this chapter, the activation mechanisms and functional significance of TRP channel's contribution to mechanotransduction is still under discussion. This is not surprising, considering that, as occurs in other cell types, diverse members of the TRP superfamily responding to mechanical energy are expressed by sensory neurons. Moreover, TRP proteins may interact forming heteromultimeric complexes. The expected functional redundancy of TRP channels participating in mechanosensation complicates the definition of the individual functional role of each channel in mechanical stimulus detection.

The identification of the molecules and/or transduction mechanisms that allow sensory neurons to differentiate stimuli of low and high intensity *in vivo* is still an important unanswered question in mechanotransduction. It is possible that non-nociceptive and nociceptive neurons express different ion channels, or the same channels but with different density.

Other factors aside from neuron sensitivity may also determine its ability to detect stimuli of different intensity, as for example a filtering of the stimulus by perireceptor structures, which could distort the efficiency of final transmission of mechanical force to the channel.

This chapter has offered a brief overview of the potential contribution of various members of the TRP channel superfamily to mechanical transduction in the somatosensory system, which leads ultimately to mechanosensation. It is evident that further work is required to fully understand the molecular and cellular processes underlying the different modalities of mechanosensation evoked by activation of the various functional classes of mechanosensitive cells, and the contribution to these processes of TRP channels.

**Acknowledgments** I thank Prof. Carlos Belmonte for helpful comments and Stuart B. Ingham for assistance with figures. I apologise for omissions of relevant work not cited due to space constraints or oversights.

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# Chapter 8

## TRP Channels in the Sensation of Heat

Chun-Hsiang Tan and Peter A. McNaughton

**Abstract** Animals must sense temperature in the external environment in order to find ambient temperatures appropriate for different activities, such as hunting or sleeping, and to avoid even brief exposure to damaging extremes of temperature. They must also sense their internal bodily temperature in order to regulate it. Some members of the TRP channel family are activated by thermal stimuli, and are consequently named the thermoTRPs; thermally sensitive ion channels are also found in the potassium and chloride channel families. When thermoTRP channels are expressed in a sensory neuron, channel opening in response to heat leads to depolarization and generation of action potentials. It is still an open question which channels are important for sensation of thermal stimuli in the external environment, and channels involved in the sensation of internal bodily temperature for the purposes of thermoregulation are even less understood. In this chapter we review the properties of thermosensitive ion channels and their roles in thermosensation and thermoregulation, with an emphasis on TRP channels.

**Keywords** TRP channels · Heat transduction · Thermosensation · Thermoregulation

### 8.1 Introduction

Thermosensation and thermoregulation are crucial for the survival of animals, which must be able both to maintain their bodily temperature within a certain range in order to carry out their physiological activities normally, and to detect and

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avoid external noxious temperatures which could cause tissue injury. Cold-blooded animals (ectotherms or poikilotherms) rely mainly on the external temperature to maintain an appropriate internal temperature, while warm-blooded animals (endotherms or homotherms) create most of their bodily heat via internal metabolic processes, and must therefore possess thermal detection mechanisms of exquisite sensitivity in order to avoid damaging hyperthermia caused by overproduction of internal heat. Animals use various strategies to achieve the goal of thermoregulation. Both ectotherms and endotherms use behavioural thermoregulation, in which sensation of the ambient temperature initiates movement towards an appropriate thermal environment. In addition to behavioural thermoregulation, mammals have a more complex thermoregulatory system under the control of the autonomic nervous system, which includes thermogenesis from brown adipose tissue (BAT) and muscle, vasoconstriction or vasodilatation of the peripheral vasculature, and evaporative heat loss via sweating. While these output mechanisms are relatively well-understood, possible mechanisms by which external and internal temperatures are sensed were mysterious until the thermally sensitive ion channels were identified and cloned. In this chapter, a brief review of the roles of heat-sensitive ion channels in mammalian thermosensation and thermoregulation will be presented, with an emphasis on TRP channels.

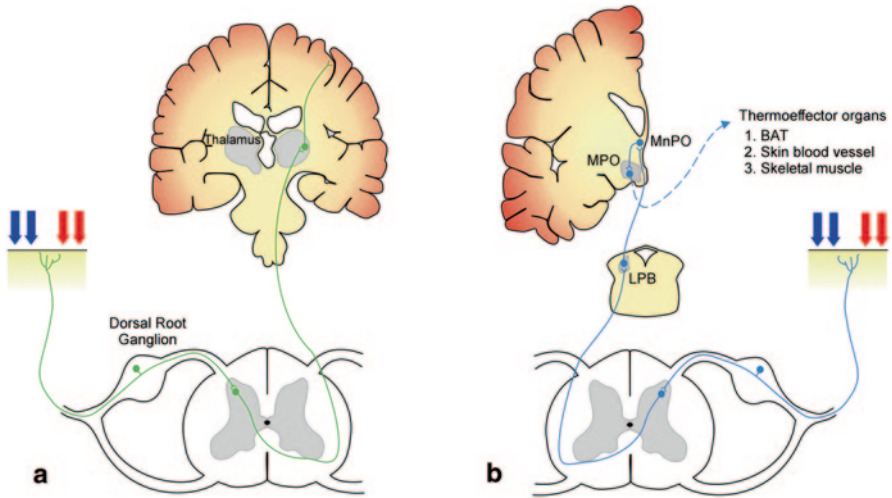
## **8.2 Thermosensation and Thermoregulation in Mammals**

### **8.2.1 Thermosensation**

Thermosensory neurons in the dorsal root ganglia (DRG) and trigeminal ganglia (TG) are pseudounipolar neurons, with a peripheral thinly myelinated or unmyelinated neurite that ends as bare dendritic terminals extending into skin, muscle, or mucosa, and a central branch that enters the spinal cord or brain stem. Thermosensitive ion channels, which depolarize the neuron when activated and thus convert thermal stimuli into firing of action potentials, are expressed in the neuronal membrane. Action potentials are transmitted to the central branch of the neuron, and from there to the spinal cord or brainstem, where thermal information is processed by excitatory and inhibitory interneurons and then transmitted onward to the thalamus and then to the primary somatosensory cortex and insular cortex to facilitate perception and discrimination of cutaneous thermosensation (Craig 2002; Todd 2010; Martinez et al. 2013). Figure 8.1a illustrates the main pathway for conscious thermosensation.

### **8.2.2 Thermoregulation**

The current concept holds that thermoregulation in mammals consists of three major components: a sensory afferent input component, a central integration/sensory



**Fig. 8.1** Pathways for thermosensation and thermoregulation. **a** Nerve terminals of somatosensory neurons in the dorsal root ganglion receive external thermal stimuli, which result in generation of action potentials. The action potentials are transmitted by the somatosensory neuron and to dorsal horn neurons in the spinal cord. Through the spinothalamic tract, the information is transmitted to the thalamus, and further relayed to the somatosensory cortex for the perception of temperature, which can lead to behavioral thermoregulation. **b** Action potentials generated in somatosensory neurons are also transmitted via dorsal horn neurons to neurons in the lateral parabrachial nucleus (LPB). LPB neurons send information to interneurons in the median preoptic (MnPO) subnucleus, which is further relayed to warm-sensitive neurons in the medial preoptic (MPO) subnucleus for integration. Preoptic warm-sensitive neurons then control thermoexecutor organs, which include brown adipose tissue (BAT), skin blood vessels, and skeletal muscles

component, and a motor efferent component. Figure 8.1b illustrates the proposed three components of thermoregulation (Morrison 2011). We discuss here the first two of these components, which are thought to contain the major sensory elements.

### 8.2.2.1 Sensory Afferent Input to Thermoregulation

There are two sensory afferent pathways important for thermoregulation. The first pathway is that used for thermosensation: DRG or TG neurons sense external warm or cold stimuli, transfer the temperature signal to secondary neurons in the spinothalamic or trigeminothalamic tract respectively, and the signal is then relayed to the cortex. Activation of the primary somatosensory cortex or insular cortex then triggers various voluntary thermoregulatory behaviors, as seen (for instance) when mice choose a preferred temperature in a thermal preference test or when humans turn on air conditioners (Flouris 2011). A good example is the initiation of warmth-seeking behavior in mice when TRPM8 is activated by menthol or cold; this behavior is reduced in TRPM8 knockout mice or by the co-application of a TRPM8 blocker (Tajino et al. 2011; Almeida et al. 2012).

A second sensory afferent pathway is initiated by the same peripheral thermosensory neurons but relays temperature signals to the pre-optic area of the hypothalamus (POAH). This pathway is crucial for controlling involuntary thermoregulation (including cold-activated vasoconstriction, sympathetic thermogenesis in BAT, and shivering thermogenesis in skeletal muscle; and heat-activated vasodilation and sweating). Cold and hot signals detected by DRG and TG neurons are transmitted through dorsal horn neurons to the external lateral subnucleus of the lateral parabrachial nucleus (LPBe) and the dorsal subnucleus of the lateral parabrachial nucleus (LPBd), respectively (Nakamura et al. 2008b; Nakamura et al. 2010). LPB neurons responsible for relaying cutaneous thermal signals then synapse on neurons in the median preoptic nucleus (MnPO), a subregion of the POAH (Nakamura et al. 2008a, b). Afferent pathways mediating cold and hot signals from the body surface to the MnPO of the POAH are essential for eliciting involuntary responses to external thermal challenges.

### 8.2.2.2 Thermoregulatory Sensorimotor Integration in the POAH

Information from the MnPO is relayed to the medial preoptic area (MPO), another subregion of the POAH (Nakamura et al. 2011). The MPO plays an important role in controlling thermoregulation by driving a potent inhibition of cold defense effector activation. The inhibitory role of the MPO on cold-defensive responses is evident from several results. First, transections of the neuraxis immediately caudal to the POAH induced a rapid and large rise in brown adipose tissue activity and increased rectal temperatures (Chen et al. 1998), and also increased cutaneous vasoconstrictor sympathetic outflow (Rathner et al. 2008). Second, reducing the activity of neurons in the MPO, either by pharmacological tools such as local injection of GABA or muscimol, a GABA<sub>A</sub> agonist, or by causing electrolytic lesions, induced cold defensive responses. Thirdly, glutamate injection, as well as electrical stimulation, suppresses cold-defensive responses and elicits vasodilatation of the paw skin and the tail (Zhang et al. 1995).

Interestingly, in the POAH, about one neuron out of five increases its rate of spontaneous discharge with an increase in POAH temperature from 32–41 °C (Nakayama et al. 1961, 1963). Moreover, local warming has the same effect as glutamate injection and electrical stimulation in suppressing cold-defensive responses and evoking cutaneous vasodilation (Kanosue et al. 1994; Zhang et al. 1995). These results indicate that the POAH neurons are intrinsically warm-sensitive and modulate their activities both according both to the local temperature of the POAH and to the afferent thermal information received from the body surface, in order to cause tonic inhibition of cold defense effector responses (Zhang et al. 1995).

However, the mechanism of how POAH neurons sense warmth remains controversial. There are two types of thermosensitive neurons in the preoptic area, primary thermosensitive neurons, which retain thermosensitivity after synaptic block, and secondary thermosensitive neurons, in which thermosensitivity disappears after synaptic block (Hori et al. 1999; Kobayashi et al. 2006). In primary thermosensitive

neurons warming increased the firing frequency by increasing the rate of depolarization between action potentials, but one group attributes this to the activation of an inward current (Hori et al. 1999; Kobayashi et al. 2006), while a second finds evidence for warmth-inhibition of an A-type potassium current (Griffin et al. 1996).

Several ion channels have been suggested to be the origin of the central thermoregulatory mechanism, such as hyperpolarization-activated cyclic nucleotide-gated channels, two-pore domain potassium channels, and the thermoTRP channels TRPV1 and TRPV4 (Wechselberger et al. 2006; Caterina 2007; Romanovsky et al. 2009). However, no thermoregulatory deficits have been reported from knockout mouse models of these channels, or from application of agonists or antagonists of hyperpolarization-activated cyclic nucleotide-gated channels or two-pore domain potassium channels. Detailed studies with TRPV1 and TRPV4 knockout mice and with TRPV1 antagonists also failed to reveal an involvement of TRPV1 and TRPV4 in warmth sensitivity in the hypothalamus (see sections below on TRPV1 and TRPV4). Several attempts with different approaches have been made to identify the thermosensitive mechanism, including a single cell transcriptome analysis of GABAergic warmth-sensitive neurons (Eberwine et al. 2011), but the mechanism remains elusive.

These studies on heat-sensitivity in the central nervous system (CNS) indicate the presence in the hypothalamus of unidentified thermosensitive ion channels which act as the thermostat of the body.

### 8.3 Heat-Sensitive Ion Channels

The use of ion channels to directly transduce information from the external environments into nerve impulses can be seen in many different modes of sensation, including thermal, acid and mechano-sensation (Waldmann et al. 1997; Coste et al. 2010). For heat-sensitivity, several families of cation channels have been reported to be heat-sensitive, including members of the TRP channel family (Vay et al. 2012; Voets 2012), the two-pore domain potassium channel family (Maingret et al. 2000; Kang et al. 2005) and the voltage-gated calcium channel family (Iftinca et al. 2006). In addition, anoctamin 1, a  $\text{Ca}^{2+}$ -activated chloride channel, has recently been reported to be heat-sensitive (Cho et al. 2012). Store-operated calcium channels of the ORAI family located in the surface membrane, while not intrinsically heat-sensitive, are modulated by the calcium-sensitive protein STIM1, in the membrane of the endoplasmic reticulum, which has also recently been reported to be heat-sensitive (Xiao et al. 2011).

Heat-sensitive ion channels expressed in the membrane of peripheral nerve terminals are responsible for transducing elevated environmental temperatures into nerve action potentials. When thermal stimuli reach the temperature threshold of the heat-sensitive ion channels, the heat-sensitive ion channels are activated, in the case of ion channels carrying an inward current, or suppressed, in the case of channels carrying an outward current, resulting in the depolarization of the neuron and the



firing of action potentials. The properties of the heat-sensitive ion channels, with emphasis on the heat-sensitive TRP channels, will be presented briefly here.

### **8.3.1 Transient Receptor Potential Channels**

The TRP channel family shares the common features of six transmembrane segments, varying degrees of sequence homology, and permeability to cations. There are 28 TRP channels in mice, while there are 27 TRP channels in humans, with TRPC2 being a pseudogene. These TRP channels can be further divided into six subfamilies, namely TRPC 1–7 (Canonical), TRPV 1–6 (Vanilloid), TRPM 1–8 (Melastatin), TRPA 1 (Ankyrin) and the more remotely related TRPP2, TRPP3, and TRPP5 (Polycystin), and TRPML 1–3 (Mucolipin) (Venkatachalam et al. 2007). TRP channels, unlike other known families of ion channels, display an enormous diversity of cation selectivities and gating mechanisms. Many TRP channels function as signal integrators, and several of them have been shown to be thermosensitive. Since all chemical and biochemical reactions are to some extent temperature-dependent, and therefore all ion channels could fall within the definition of temperature-sensitive, we define thermosensitivity from the  $Q_{10}$  value, which is calculated as the relative increase in current amplitude at a specific voltage when the temperature increases by 10°C. Here, a brief overview of the heat-sensitive TRP channels which have  $Q_{10}$  values higher than 5 will be presented. These are TRPV1, TRPV2, TRPV3, TRPV4, TRPM2, TRPM3, TRPM4 and TRPM5 (Voets 2012).

#### **8.3.1.1 Transient Receptor Potential Vanilloid 1 (TRPV1)**

The activation of ion channels by noxious heat in somatosensory neurons was first characterized in 1996 (Cesare et al. 1996). The ion channel responsible for heat sensitivity in at least some primary afferent neurons was cloned in the following year (Caterina et al. 1997), and was later named TRPV1 to emphasize its membership of the TRP ion channel family. TRPV1 is a nonselective cation channel displaying outward rectification that can be activated by temperature >42°C with a  $Q_{10}$  value of 25 (Vlachova et al. 2003) and  $P_{Ca}/P_{Na}$  value of around 10 (Owsianik et al. 2006). TRPV1 can also be activated by a wide range of agonists, including capsaicin, resiniferatoxin and extracellular protons, and is inhibited by capsazepine, N-(4-tertiarybutylphenyl)-4-(3-chloropyridin-2-yl)tetrahydropyrazine-1(2H)-carboxamide (BCTC) and ruthenium red, amongst others. TRPV1 was found to be mainly expressed in C-fibers and A $\delta$  nociceptive fibers of somatosensory neurons that express the neuropeptides substance P, neurokinin A, and CGRP, indicating its role as a sensor of noxious heat stimuli (Tominaga et al. 1998; Kobayashi et al. 2005; Julius 2013).

TRPV1 KO mice show a partial deficit in sensing noxious heat, but they still withdraw their tails from hot water (Caterina et al. 2000), and TRPV1 KO mice

also show no difference from wild type mice in two-plate temperature preference tests (Pogorzala et al. 2013). In addition, no differences in heat responses could be observed in polymodal nociceptors between WT and TRPV1 KO mice, and the role of TRPV1 seems to be restricted to the C-heat fiber class (Woodbury et al. 2004; Lawson et al. 2008). These findings demonstrate the existence of additional TRPV1-independent mechanisms of heat sensation.

Several studies have sought to determine whether TRPV1 is involved in thermoregulation, in pursuit of the long-sought central mechanism responsible for regulation of body temperature. A possible role of TRPV1 in thermoregulation was supported by evidence showing that TRPV1 blockade with a highly selective TRPV1 antagonist, AMG 517, elicits a marked and plasma concentration-dependent hyperthermia in rodents and humans (Gavva et al. 2007, 2008). However, a peripherally restricted TRPV1 antagonist causes similar increases in body core temperature, demonstrating that the site of action for TRPV1 blockade is outside the blood-brain barrier, and that TRPV1 is therefore not directly involved in central thermoregulation in the hypothalamus (Tamayo et al. 2008). In addition, several studies have failed to reveal any difference between TRPV1 KO and WT mice in basal deep body temperature or in the change in the deep body temperature in responses to thermal challenges (Szelenyi et al. 2004; Iida et al. 2005; Garami et al. 2011). All these results suggest that TRPV1 is unlikely to be a thermosensitive mechanism playing an important role in central thermoregulation.

### 8.3.1.2 Transient Receptor Potential Vanilloid 2 (TRPV2)

Two years after the cloning of TRPV1, a TRPV1 homologue with a high threshold for noxious heat was discovered, and was subsequently named TRPV2 (Caterina et al. 1999). TRPV2 carries an outwardly rectifying nonselective cation current with  $P_{Ca}/P_{Na}$  of around 3 (Clapham et al. 2005; Owsianik et al. 2006). TRPV2 was once thought to be a candidate for the TRPV1-independent mechanism for heat sensation, because the channel has an even higher temperature threshold than TRPV1 ( $\sim 52^{\circ}\text{C}$ ) with a  $Q_{10}$  value higher than 100 (Leffler et al. 2007), and is expressed in medium- to large- diameter myelinated sensory neurons (Caterina et al. 1999). TRPV2 is also activated by hypo-osmolarity, membrane stretch and cannabinoids in addition to heat (Muraki et al. 2003; Qin et al. 2008), and is inhibited by ruthenium red, lanthanum, and SKF96365 (Muraki et al. 2003; Clapham et al. 2005; Qin et al. 2008; Zhang et al. 2012a). The activation of TRPV2 by 2-APB is controversial, with some reports showing that TRPV2 can be activated by 2-APB (Hu et al. 2004; Neeper et al. 2007), and others not (Chung et al. 2004; Bender et al. 2005). In addition to expression in somatosensory neurons, TRPV2 is also expressed in the epithelium of several different tissues (including pancreatic duct, mammary gland, parotid gland, submandibular gland, renal tubule and tracheal gland) and in immune cells (Kowase et al. 2002; Link et al. 2010).

The wide distribution of expression in various tissues suggests a possible range of physiological functions mediated by TRPV2 but the physiological or pathologi-

cal functions mediated by TRPV2 still remain unclear, because behavioral assays of TRPV1/TRPV2 double knockout mice, or TRPV2 knockout mice treated with resiniferatoxin to desensitize TRPV1-expressing afferents, reveal no thermosensory deficits as a consequence of absence of TRPV2 (Park et al. 2011). Whether TRPV2 could be involved in thermoregulation remains unclear. However, the finding showing that the overall pattern and amplitude of the oscillation in core body temperature was not different in TRPV2 knockout mice from WT mice, along with the high activation threshold by heat, makes the possibility less likely (Park 2008).

### 8.3.1.3 Transient Receptor Potential Vanilloid 3 (TRPV3)

TRPV3, also a homologue of TRPV1, was identified by searching for sequences with a high degree of similarity to TRPV1 in the human genome (Peier et al. 2002; Smith et al. 2002) or by searching for TRP-related expressed sequence tags (ESTs) in a human testis library (Xu et al. 2002). TRPV3 carries an outwardly rectifying nonselective cation current with a  $P_{Ca}/P_{Na}$  value of around 12 (Xu et al. 2002; Clapham et al. 2005; Owsianik et al. 2006). TRPV3 can be activated by warmth with an activation threshold at a temperature around 31 °C and has a  $Q_{10}$  value higher than 6 (Peier et al. 2002; Xu et al. 2002). In addition to temperature, TRPV3 can also be activated by carvacrol, eugenol and thymol, which are found in plants such as oregano, savory, clove and thyme (Xu et al. 2006b). TRPV3 is also activated by 2-APB and is inhibited by ruthenium red (Hu et al. 2004; Clapham et al. 2005). In addition, TRPV3 exhibits a unique property known as sensitization, which causes progressive increases in its response with repeated application of a stimulus (Peier et al. 2002).

TRPV3 is mainly expressed in skin keratinocytes and has been proposed to act as a thermosensor, the excitation of which could be transferred to primary afferent terminals of sensory neurons by an unknown mediator (Peier et al. 2002). TRPV3 null mice on an intercrossed C57BL6/129 J background were initially reported to have some relatively minor deficits in responses to innocuous and noxious heat but not in other sensory modalities (Moqrich et al. 2005). However, TRPV3 knockout mice on a homogeneous C57BL6 background exhibited no obvious alterations in thermal preference behaviour, while the absence of TRPV3 on a 129S6 background resulted in a more restricted range of occupancy centered around cooler floor temperatures (Huang et al. 2011). In addition, TRPV3 knockout mice showed no deficits in acute heat nociception on either a C57BL6 or a 129S6 background (Huang et al. 2011). These results suggest that TRPV3 is not a major contributor for thermosensation, and the previously reported deficits in responses to innocuous and noxious heat in the TRPV3 null mice on the intercrossed C57BL6/129J background are possibly due to inhomogeneous inheritance of non-TRPV3 determinants. The warmth-sensitive TRPV3 could plausibly be involved in thermoregulation, but the finding that mice in which TRPV3 has been genetically inactivated appeared morphologically normal and have a normal internal temperature does not favour this idea (Moqrich et al. 2005).

### 8.3.1.4 Transient Receptor Potential Vanilloid 4 (TRPV4)

TRPV4 was cloned with an initial motivation to find osmotically-sensitive channels (Liedtke et al. 2000; Strotmann et al. 2000). It was almost 2 years after the cloning of TRPV4 that it was found to be thermo-sensitive (Guler et al. 2002). TRPV4 carries an outwardly rectifying nonselective cation current that can be activated by non-noxious heat at a temperature higher than 25 °C (Clapham 2003) with a  $Q_{10}$  value of around 10 (Guler et al. 2002) and  $P_{Ca}/P_{Na}$  value of around 9.3 (Strotmann et al. 2003). In addition to warmth and hypotonic stress, TRPV4 can also be activated by the compound bisandrographolide from the Chinese herbal plant *Andrographis paniculata* and by a specific TRPV4 agonist, GSK1016790A (Smith et al. 2006; Willette et al. 2008). TRPV4 is also inhibited by ruthenium red and by a specific TRPV4 antagonist, HC-06704 (Clapham et al. 2005; Everaerts et al. 2010). The expression of TRPV4 has been reported in a variety of tissues, including kidney, corneal epithelial cells (Pan et al. 2008), cerebral microvascular endothelial cells (Ma et al. 2008), cortical astrocytes (Benfenati et al. 2007), tracheal epithelial cells (Lorenzo et al. 2008), and keratinocytes (Guler et al. 2002).

Like TRPV3, the fact that TRPV4 is expressed in keratinocytes led to the proposal that activation of TRPV4 in keratinocytes could be transferred to sensory neurons via an unidentified extracellular mediator. However, as noted above, mice deficient in both TRPV3 and TRPV4 show thermal preference behavior similar to wild type mice on a thermal gradient and little or no change in acute heat perception (Huang et al. 2011). The findings suggest that TRPV4 probably also makes a very limited contribution to heat sensation.

TRPV4 also seems not to be involved in thermoregulation. Application of a TRPV4 antagonist did not affect core body temperature (Everaerts et al. 2010). Second, no difference in basal body temperature recordings and no difference in core body temperature after cold stress or heat stress was found between TRPV4 KO and WT mice (Liedtke et al. 2003; Lee et al. 2005).

### 8.3.1.5 Transient Receptor Potential Melastatin 2 (TRPM2)

TRPM2 was cloned in 1998 with an initial goal of finding the genes responsible for several diseases, such as autoimmune polyglandular disease type I, bipolar affective disorder, and nonsyndromic hereditary deafness, and was confirmed to be located at 21q22.3 (Nagamine et al. 1998). Of these diseases, bipolar disorder was found to be associated with TRPM2 (Yoon et al. 2001; Xu et al. 2006a). TRPM2 carries a voltage-insensitive nonselective cation current that was found later to be activated by heat at a temperature higher than 35 °C with  $Q_{10}$  value at around 15.6 (Togashi et al. 2006) and  $P_{Ca}/P_{Na}$  value of 0.5–1.6 (Perraud et al. 2001). In addition to heat, the activity of TRPM2 can also be synergistically modulated by intracellular ADP-ribose (ADPr) and  $Ca^{2+}$  (Heiner et al. 2006). TRPM2 can also be activated by reactive oxygen species (ROS), such as  $H_2O_2$  (Hara et al. 2002; Wehage et al. 2002), and enhanced by additional factors, such as cyclic ADP-ribose and nicotinic

acid adenine dinucleotide phosphate (NAADP) (Kolisek et al. 2005; Beck et al. 2006). TRPM2 is inhibited by permeating protons (both intracellular and extracellular acidic pH) (Du et al. 2009; Starkus et al. 2010) and intracellular adenosine monophosphate (AMP) (Kolisek et al. 2005; Beck et al. 2006).

TRPM2 is highly expressed in brain (with the vast majority in non-neuronal microglial cells) and in immune cells such as macrophages and neutrophils (Naziroglu 2011), and has also been reported to be expressed in DRG neurons (Naziroglu et al. 2014). However, TRPM2 KO mice showed no deficits in noxious heat sensitivity when tested with the Hargreaves test and hot plate test at 52 and 55°C (Haraguchi et al. 2012). Thermal hyperalgesia was, however, found to be attenuated in carrageenan-induced inflammatory pain and sciatic nerve injury-induced neuropathic pain models (Haraguchi et al. 2012). It was proposed that rather than being a direct effect mediated via TRPM2 expressed in sensory nerve terminals, the effect of TRPM2 deletion on pain was due to TRPM2 expressed in macrophages and microglia aggravating peripheral and spinal pronociceptive inflammatory responses and hence contributing to the pathogenesis of inflammatory and neuropathic pain (Haraguchi et al. 2012).

TRPM2 KO mice have a similar body temperature to WT mice at ambient temperatures (Uchida et al. 2011). However, the activation threshold of TRPM2 at around body temperature (Togashi et al. 2006) makes TRPM2 a possible candidate as the long-sought thermosensitive mechanism for thermoregulation. In addition, TRPM2 KO mice have been reported to have higher energy expenditure (Zhang et al. 2012b), which in general is correlated with thermoregulation. More detailed studies into the thermoregulatory phenotype of TRPM2 KO mice are needed to determine whether TRPM2 may be involved in thermoregulation.

### 8.3.1.6 Transient Receptor Potential Melastatin 3 (TRPM3)

TRPM3 has the largest number of splice variants within the TRP channel family (11 splice variants have been reported) (Oberwinkler et al. 2007; Thiel et al. 2013) and was first identified by sequencing projects. TRPM3 carries an outwardly rectifying non-selective cation current that can be activated by heat (>40°C) with a  $Q_{10}$  value of 7.2 (Vriens et al. 2011) and  $P_{Ca}/P_{Na}$  at 0.1–10 (depending on splice variant) (Owsianik et al. 2006). TRPM3 exhibits constitutive activity when overexpressed in HEK293 cells. Agonists of TRPM3 include pregnenolone sulphate, nifedipine, and  $\beta$ -cyclodextrin (Wagner et al. 2008; Naylor et al. 2010). Secondary metabolites from citrus fruit flavanones, such as naringenin (the predominant flavanone in grapefruits), hesperetin (the predominant flavanone in oranges), and fabecea (from the pea and bean family) selectively inhibit TRPM3 channel activation (Felgines et al. 2000; Erlund et al. 2001; Straub et al. 2013a, b). TRPM3 channels are prominently expressed in various tissues, including kidney, liver, ovary, brain, spinal cord, pituitary, vascular smooth muscle, and testis (Grimm et al. 2003; Lee et al. 2003; Naylor et al. 2010) and are also found to be expressed in small diameter DRG neurons (Vriens et al. 2011).

TRPM3 knockout mice exhibited reduced but not absent sensitivity to noxious heat, similar to what was seen with TRPV1 knockout mice. This result is corroborated by the effect of block or deletion of TRPV1 and TRPM3 in sensory neurons; up to 50% of DRG neurons from TRPM3 knockout mice are still heat-responsive after treatment with AMG9810, a selective TRPV1 antagonist (Vriens et al. 2011). These results demonstrate that both TRPV1 and TRPM3 contribute to the sensation of noxious heat, but that there are additional novel heat-sensitive mechanisms independent of TRPV1 and TRPM3.

Whether TRPM3 is involved in thermoregulation in TRPM3 KO mice is undetermined, because no detailed studies have been reported. The only result available is that the body temperatures of TRPM3 KO mice are not statistically different from WT mice (Vriens et al. 2011).

### 8.3.1.7 Transient Receptor Potential Melastatin 4 (TRPM4)

Within the TRPM channel family most channels are non-selective cationic channels, but TRPM4 and TRPM5 are not permeable to divalent cations (Guinamard et al. 2010). TRPM4 carries an outwardly rectifying current that has a  $Q_{10}$  value of around 8.5 between 15 and 25 °C (Talavera et al. 2005). In addition to heat, TRPM4 can be activated directly by intracellular calcium with an  $EC_{50}$  of 20  $\mu$ M (Ullrich et al. 2005). Decavanadate and BTP2 are also activators of TRPM4 (Nilius et al. 2004a; Takezawa et al. 2006). Intracellular ATP can inhibit TRPM4 with an  $IC_{50}$  value of 1.7  $\mu$ M (Nilius et al. 2004b). Other reported inhibitors of TRPM4 include flufenamic acid, 9-phenanthrol, MPB-104, glibenclamide, and clotrimazole (Abriel et al. 2012). TRPM4 expression is widespread and, in humans, it has been detected at high levels in the heart, pancreas, placenta, and prostate and at lower levels in the kidney, skeletal muscle, liver, intestines, thymus, and spleen (Abriel et al. 2012).

TRPM4 mRNA has also been shown to be expressed in trigeminal ganglion and in DRG at all the anatomical segments of the spinal cord tested (Jang et al. 2012; Vandewauw et al. 2013). However, no studies focusing on a possible thermosensory or thermoregulatory phenotype of TRPM4 KO mice have been reported. A possible role for TRPM4 in thermosensation and thermoregulation needs further investigation.

### 8.3.1.8 Transient Receptor Potential Melastatin 5 (TRPM5)

TRPM5, like TRPM4, is impermeable to divalent cations (Vandewauw et al. 2013). TRPM5 carries an outwardly rectifying current that has a  $Q_{10}$  value of around 10.3 between 15 and 25 °C (Talavera et al. 2005). In addition to warmth, TRPM5 can also be activated directly by intracellular calcium at an even lower concentration than TRPM4, with  $EC_{50}$  at 0.7  $\mu$ M, whereas higher concentrations are inhibitory (Prawitt et al. 2003; Ullrich et al. 2005). TRPM5 can be inhibited by triphenylphosphine oxide (Palmer et al. 2010). External acidification can also cause a rapid and



reversible block of the current mediated by TRPM5, with  $IC_{50}$  pH of 6.2 and a slower irreversible enhancement of current inactivation (Liu et al. 2005). TRPM5 was first cloned from mammalian taste buds (Perez et al. 2002) and has been shown to confer thermal sensitivity to sweet taste (Damak et al. 2006; Liman 2007) (See Chap. 6 by Talavera in this book). A temperature increase from 15 to 35 °C significantly enhances the response of the gustatory nerve to sweet compounds in WT but not in TRPM5 KO mice (Talavera et al. 2005). TRPM5 expression was later found to be widespread and includes chemosensory organs (the main olfactory epithelium and the vomeronasal organ) (See Chap. 5 by Restrepo et al. in this book) and epithelial cells of the respiratory and gastrointestinal tract (Kaske et al. 2007).

TRPM5 mRNA has also been reported to be detectable in DRG at all anatomical segments of the spinal cord tested (Vandewauw et al. 2013). However, no studies of the role of TRPM5 in either thermosensation or thermoregulation have been reported, and a possible role of TRPM5 in these physiological functions is undetermined.

### 8.3.2 Two-Pore Domain Potassium Channels

Mammalian two-pore domain potassium channels (K2P) are encoded by 15 known genes (KCNK family). The two-pore domain potassium channels share the common features of four transmembrane segments, two pore domains, an extended M1P1 extracellular loop and intracellular amino (N) and carboxyl (C) termini. However, they have low sequence identity outside the pore regions. The K2P channels can be distributed into six subfamilies [two-pore domain weak inwardly rectifying  $K^+$  channel (TWIK), TWIK-related  $K^+$  channel (TREK), TWIK-related acid-sensitive  $K^+$  channel (TASK), TWIK-related alkaline-activated  $K^+$  channel (TALK), TWIK-related halothane-inhibited  $K^+$  channel (THIK), and TWIK-related spinal cord  $K^+$  channel (TRESK)]. These two-pore domain potassium channels give rise to leak  $K^+$  currents, and have also been shown to be important in determining neuronal excitability and cell firing (Enyedi et al. 2010). Among the two-pore potassium channels, TREK-1, TREK-2, and TRAAK have been shown to be thermo-sensitive (Maingret et al. 2000; Kang et al. 2005).

TREK-1 is activated by heat with a  $Q_{10}$  from 22 to 32 °C of ~7-fold, while a temperature decrease from 22 to 16 °C inhibits TREK-1 basal activity. Noteworthy is that cell integrity is required for temperature sensitivity, because TREK-1 in the inside-out state is not sensitive to heat (Maingret et al. 2000). In addition to the sensitivity to temperature, TREK-1 can be activated by arachidonic acid, unsaturated fatty acids, lysophospholipids, volatile anesthetics, mechanical stress, and internal acidification, and is inhibited by barium and quinidine (Goldstein et al. 2005; Ma et al. 2011).

TREK-2 can also be activated by heat, with a ~20-fold increase in current amplitude from 24 to 42 °C. Cell integrity is also required for temperature sensitivity, suggesting the participation of an intracellular second messenger in temperature transduction (Kang et al. 2005). TREK2 is activated by arachidonic acid, docosa-



hexaenoic acid, chloroform, halothane, and isoflurane, mechanical stress, intracellular acidification and riluzole, and is inhibited by quinidine (Lesage et al. 2000; Goldstein et al. 2005).

TRAAK has a similar sensitivity to heat to TREK-2, with a ~20-fold increase in current amplitude with a temperature increase from 24 to 42 °C, and like TREK-1 and TREK-2, cell integrity is required for thermosensitivity (Kang et al. 2005). Arachidonic acid, mechanical stress, unsaturated fatty acids, lysopholipids and riluzole can activate TRAAK in addition to heat, and gadolinium can inhibit TRAAK (Goldstein et al. 2005).

TREK-1, TREK-2 and TRAAK have been shown to be expressed in both somatosensory neurons and the CNS (Maingret et al. 2000; Medhurst et al. 2001; Pollema-Mays et al. 2013) and therefore could be involved in thermosensation and/or thermoregulation. TREK-1 and TRAAK KO mice were reported to be more sensitive to heat stimuli, but not to cold (Alloui et al. 2006)(Noel et al. 2009). Interestingly, in TREK-1/TRAAK double knockout mice, heat hypersensitivity persists and cold hypersensitivity develops. Recently, TREK-2 KO mice were shown to have increased excitability to thermal stimuli above 40 °C and TREK-1/TREK-2/TRAAK triple knockout mice also express hypersensitivity to heat (Pereira et al. 2014). These results suggest that inhibition of these K2P channels by cold could promote firing in cold-sensitive neurons, while in response to heat, activation of these channels could act to dampen the effect of activation of TRP channels and so to reduce heat sensitivity.

It remains unknown whether two-pore domain potassium channels are involved in thermoregulation. None of the two-pore domain potassium channel KO animals have been found to have any obvious deficits in thermoregulation, but there is a lack of definitive studies and participation in thermoregulation remains a possibility.

### **8.3.3 Calcium-Activated Chloride Channel Anoctamin 1 (ANO1)**

The calcium-activated chloride channel ANO1 was recently reported to be heat-sensitive. ANO1 is activated by heat (>44 °C) with a  $Q_{10}$  value of 19.4 (Cho et al. 2012) and by intracellular calcium. ANO1 can also be activated by other divalent cations, including  $Ba^{2+}$ ,  $Sr^{2+}$ , and  $Ni^{2+}$ . Niflumic acid, often considered a specific blocker for calcium-activated chloride channels, blocks ANO1. Other blockers of ANO1 have been investigated and include T16inh-A01, CaCCinh-A01, tannic acid, eugenol, and benzbromarone (Pedemonte et al. 2014).

ANO1 has been shown to be expressed in epithelial cells of pulmonary bronchioles, pancreatic acinar cells, retinal cell layers, proximal renal tubules, submandibular glands, Leydig cells, and sensory neurons of dorsal root ganglia (Yang et al. 2008). ANO1 knockdown mice and ANO1 conditional knockout mice were both shown to have deficits in sensing noxious heat, showing that ANO1 is involved in heat sensation (Cho et al. 2012). However, there are no reports showing whether ANO1 may be involved in thermoregulation.

## 8.4 Concluding Remarks

In the last two decades several TRP channels have been shown to be activated by heat, and both TRPV1 and TRPM3 have been demonstrated to play a role in heat sensation *in vivo*. Thermal sensitivity is not confined to the TRP channel family, however, as members of the two-pore domain potassium channel family and the calcium-activated chloride channel anoctamin 1 are also involved in the detection of heat from the external environment. The evidence for an involvement of other heat-sensitive TRP channels in thermosensation *in vivo* is not currently persuasive. However, it is still undetermined whether the combination of these known heat-sensitive mechanisms is enough to fully explain the exquisite sensitivity of animals to thermal stimuli, or whether there are other novel thermosensitive mechanisms remaining to be discovered. More importantly, none of the existing knockout animals show significant deficits in thermoregulation, and an understanding of the molecular mechanisms underlying the regulation of the core temperature of mammals remains elusive.

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# Chapter 9

## TRP Channels in Cold Transduction

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María Pertusa and Rodolfo Madrid

**Abstract** In the somatosensory system, cold thermoreceptor neurons and cold nociceptors are responsible for the detection of environmental low temperatures. The underlying machinery is far from simple; it is a result of the participation of several classes of transduction and voltage-gated ion channels that functionally coexist to give shape to the cold-induced receptor potential and subsequent action potential firing in response to cold stimulation. The cold-induced electrical responses begin in the free nerve endings of these sensory neurons, where a subgroup of thermo-sensitive **Transient Receptor Potential** channels (thermoTRPs) plays a critical role. These channels have evolved as molecular thermal sensors activated by a wide range of cold temperatures, and they have been proposed as key elements of the transduction machinery responsible for detection of environmental cold in primary somatosensory neurons. In this chapter, we summarize the most important functional properties of the primary sensory neurons involved in cutaneous cold detection, and the corresponding role of the thermoTRP channels TRPM8, TRPA1 and TRPC5 in cold transduction.

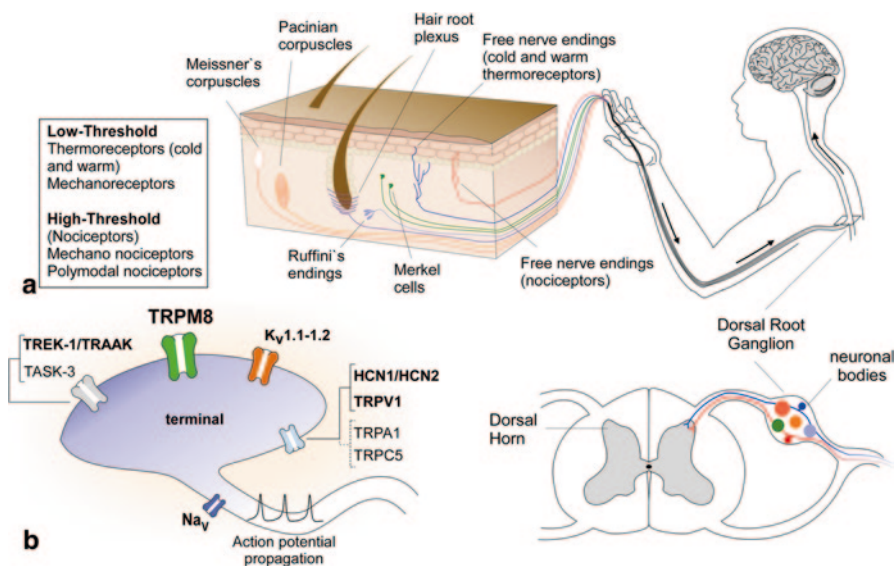
**Keywords** Cold transduction · TRPM8 · TRPA1 · TRPC5 · Primary sensory neurons · TRP channels

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## 9.1 Introduction

Primary sensory neurons of the peripheral somatosensory system are responsible for converting a wide range of environmental physical and chemical stimuli into electrical signals. Sensory signals begin at the nerve endings of these pseudounipolar neurons that innervate the skin and exposed mucosae, which home the molecular machinery necessary for the detection and transduction of these stimuli into a receptor potential (Fig. 9.1). The soma of primary somatosensory neurons is located in the dorsal root ganglia (DRG) and trigeminal ganglia (TG). These neurons project one axonal branch peripherally innervating the skin and mucosae, and the other branch centrally to dorsal horn neurons at the spinal cord and brain stem nuclei,



**Fig. 9.1** Sensory innervation of the skin. **a.** Simplified schematic representation of the different primary sensory neurons innervating the skin. **b.** In the dorsal root ganglion at right, the somas of the different subclasses of primary sensory neurons that project to dorsal horn are represented in different colors. The nerve ending of a cold thermoreceptor neuron (CTN) is shown in the *left panel*. The site of cold transduction (terminal) is depicted as separated from the site of action potential generation and propagation. In this scheme, TRPM8 is the most prominent channel, representing its contribution to cold sensing. Kv1.1-1.2 channels (the molecular counterpart of the break potassium current  $I_{KD}$ ), TREK-1/TRAAK and TASK-3 background potassium channels are also important components of the molecular machinery underlying cold temperature transduction. HCN1 and, to a lower extent, HCN2 channels (the molecular counterpart of  $I_h$  current) contribute to give shape to the net electrical response to temperature reductions in primary somatosensory neurons. The heat- and capsaicin-activated TRPV1 channel is also expressed in a large subpopulation of CTNs (~25%); TRPA1 is only expressed in a small subpopulation of CTNs (~5% of the TRPM8-expressing neurons). Although depicted, further studies are necessary to determine the role of TRPC5 in cold transduction. Among  $Na_v$  channels,  $Na_{v1.8}$  channels are critical to allow impulse generation in primary sensory neurons at very low temperatures

where the information is transmitted towards the central integrative areas of the brain.

TG and DRG comprise a variety of neurons responding to a wide range of mechanical stimuli, chemical agents of different nature, and temperature, and the activation of functional subpopulations of the primary sensory neurons evoke diverse sensations of touch, itch, irritation, cold, heat and pain. Thus, primary somatosensory receptors can be divided into cold and warm thermoreceptors, mechanoreceptors and nociceptors. Based on the intensity of the stimuli necessary to initiate a sensory signal, primary somatosensory neurons can be separated into low-threshold and high-threshold receptors. Under this general classification, low-threshold receptor neurons correspond to thermoreceptors (cold and warm) and low-threshold mechanoreceptors, whereas high-threshold receptors correspond to nociceptors, which include polymodal nociceptors and mechanonociceptors (Fig. 9.1a).

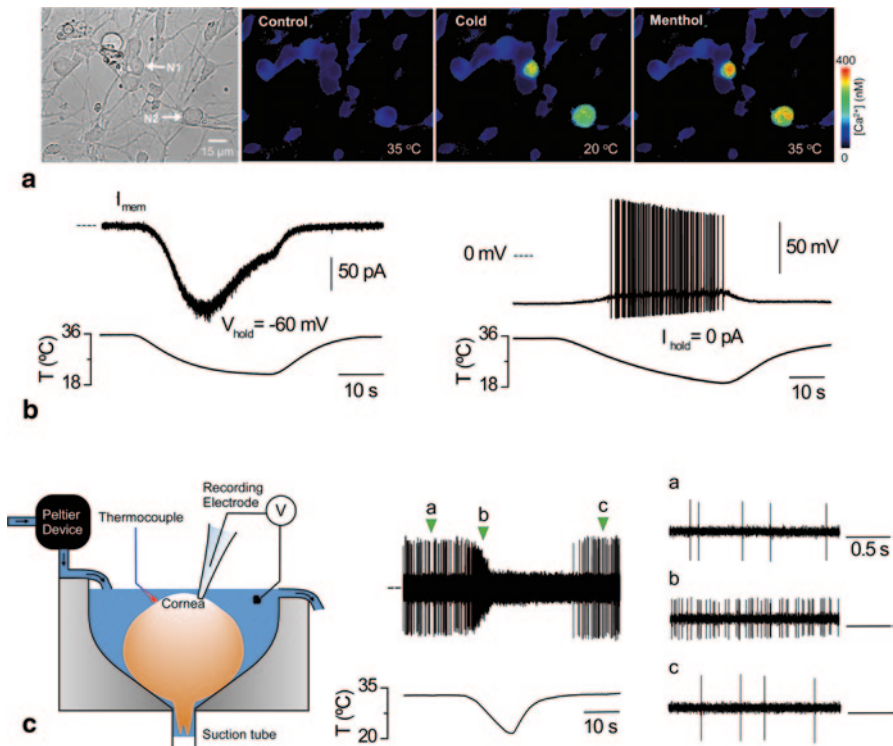
The wide diversity of cold sensations that arise from the exposure of the skin and mucosae to environmental temperature reductions varies from comfortable cold to intense pain, depending on the intensity of the thermal stimulus. The perception of cold as an innocuous temperature drop takes place when the surface of the skin is cooled by as little as 1 °C, or even less, and cold normally feels painful or noxious at temperatures below 15 °C. The detection of innocuous and noxious cold is mediated by cold thermoreceptors and nociceptors respectively, expressing a large variety of transduction and voltage-gated channels. These channels work concertedly in generating the cold-induced receptor potential and subsequent action potential firing in response to an environmental temperature decrease. In the last years, the identification of ion channels with a strong thermal sensitivity, including several cold- and heat-activated members of the superfamily of **Transient Receptor Potential (TRP)** (also called thermoTRP channels), has shed light on the molecular mechanisms underlying the diverse sensory modalities that emerge from the activation of distinct sets of peripheral receptors, including cold-sensitive neurons (Vriens et al. 2014).

## 9.2 Cold Thermoreceptor and Cold Nociceptor Neurons

In mammals, cold thermoreceptor neurons (CTNs) correspond to subpopulations of small diameter unmyelinated primary afferent C-fibers and medium diameter thinly myelinated A $\delta$ -fibers, in a proportion that varies among different species and somatic territories (Hensel and Zotterman 1951b; Iriuchijima and Zotterman 1960; Hensel 1981; Heppelmann et al. 1990). CTNs show a wide range of temperature thresholds, and can be separated into low-threshold and high-threshold cold-sensitive neurons, the latter having a role in the detection of cold discomfort under physiological conditions (Belmonte et al. 2009). At a normal skin temperature of 33 °C, cold thermoreceptors detecting and encoding innocuous or moderate cold typically exhibit spontaneous electrical activity, mainly in a beating (regular) pattern of action potential firing. The firing of these exquisitely temperature-sensitive neurons increases in response to temperature reductions of less than 1 °C, and is reduced

by heating. These cold afferents are sensitized by micromolar concentrations of the natural cold-mimetic compound menthol (Hensel and Zotterman 1951a; Brock et al. 2001). Cold thermoreceptor neurons comprise about 8–15% of the neuronal population in DRG and TG.

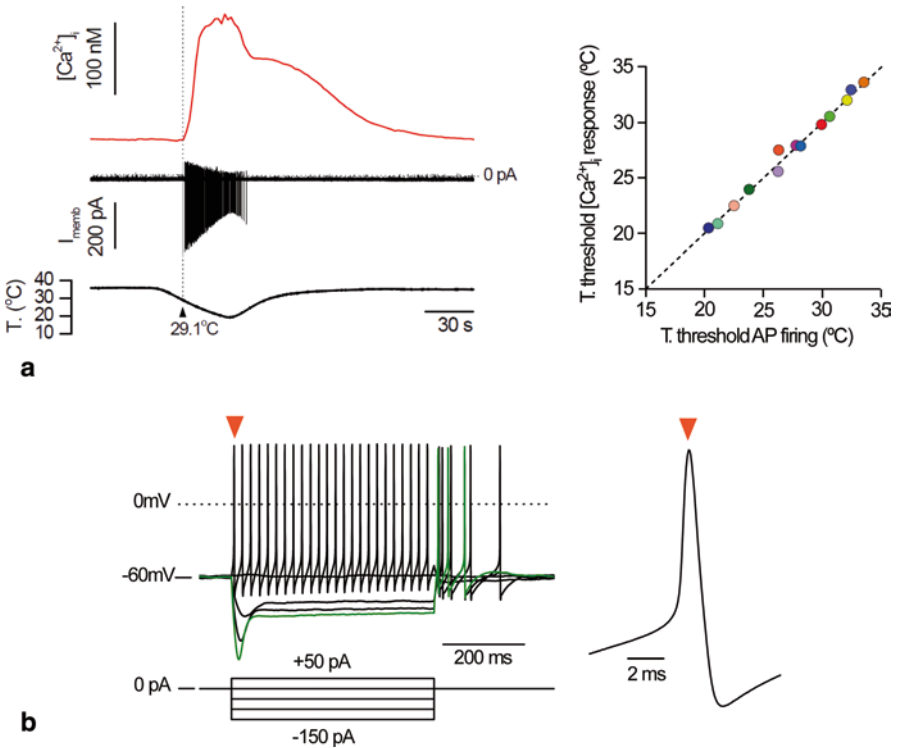
The functional properties of primary somatosensory neurons that respond to innocuous and moderate cold have been extensively characterized by  $\text{Ca}^{2+}$ -imaging, patch-clamping and extracellular recordings in several somatic territories (Fig. 9.2). In culture, these neurons respond to cooling with a depolarizing inward current. The depolarization induces an increase in the action potential firing followed by a rise in intracellular  $\text{Ca}^{2+}$  concentration that depends on external  $\text{Ca}^{2+}$ . In CTNs,



**Fig. 9.2** Calcium imaging and electrophysiological tools in the study of thermal responses in cold thermoreceptor neurons. **a.** Transmitted and pseudocolor ratiometric  $[\text{Ca}^{2+}]_i$  images cultured trigeminal neurons showing the effects of cold and menthol on its intracellular calcium concentration. Note that the same neurons are sensitive to both stimuli (Madrid and Viana, unpublished). Scale bar in *left* panel, 15 μm. **b.** *Left*, simultaneous recording of membrane current using patch clamp technique (*top* trace) and bath temperature (*bottom* trace) during a cooling ramp from 34 to 22°C in a cold-sensitive trigeminal neuron ( $V_{\text{hold}} = -60$  mV). *Right*, simultaneous recording of membrane potential (*top* trace) and bath temperature during a cooling ramp in a cold-sensitive neuron recorded in current-clamp mode ( $I_{\text{hold}} = 0$  pA) (González and Madrid, unpublished). **c.** Extracellular recording of nerve terminal impulses (NTI) activity in a corneal cold-sensitive neuron in response to cooling (González and Madrid, unpublished; Scheme at *left* modified from Parra et al. 2010)

intracellular  $\text{Ca}^{2+}$  increases in response to cold stimulation are mainly due to the activation of voltage-gated  $\text{Ca}^{2+}$  channels during action potential firing. As a consequence, there is a tight correlation between temperature threshold for firing of action potentials and for  $[\text{Ca}^{2+}]_i$  elevations in each individual neuron (Viana et al. 2002; Madrid et al. 2006) (Fig. 9.3a). Canonical CTNs fire short duration action potentials ( $\sim 1$  ms at the half-amplitude of the depolarizing phase) (Fig. 9.3b) and are characterized by a low rheobase current and by the presence of a prominent hyperpolarization-activated current ( $I_h$ ), mediated by HCN1 and, to a lower extent, HCN2 channels (Reid et al. 2002; Viana et al. 2002; Madrid et al. 2006, 2009; Orío et al. 2009, 2012).

As mentioned before, the vast majority of cold thermoreceptor neurons are sensitive to menthol, a natural cyclic terpene alcohol that induces cold sensations

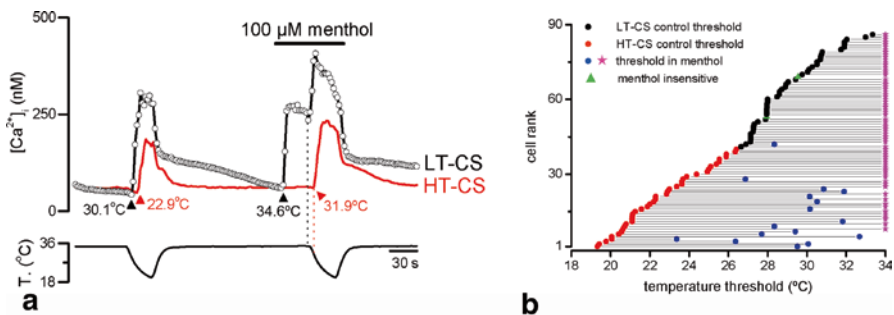


**Fig. 9.3** The intracellular calcium increase in response to cold in cultured cold-sensitive neurons depends on the action potential firing. **a** *Left*, simultaneous recording of  $[\text{Ca}^{2+}]_i$  signals (*top trace*), action currents (*middle*) and bath temperature (*bottom*) during a cooling ramp in a cold-sensitive neuron in culture, recorded in cell-attached condition. *Right*, scatter plot of thresholds for action potential firing and  $[\text{Ca}^{2+}]_i$  signals in response to cooling in 14 cold-sensitive neurons (each neuron has been color coded). The dotted line represents the unity line (González and Madrid, unpublished). **b** *Left*, typical voltage responses (*upper traces*) to 500 ms hyperpolarizing and depolarizing current pulses (*lower traces*) of a trigeminal CTN. Note the fast repetitive discharge, the strong sag and the presence of rebound firing (*in green*). The first action potential of the train (*orange arrow*) is shown at *right* (González and Madrid, unpublished)

(Hensel and Zotterman 1951a). Menthol sensitivity in cold-sensitive fibers (and cold-sensitive cultured neurons) is manifested as an increase in the action potential firing frequency in response to this so-called cooling compound at 33 °C. Expressed differently, menthol shifts the temperature threshold of the cold-induced response to higher temperatures. This is not a trivial point, since high-threshold cold-sensitive neurons that do not respond to menthol at 33 °C could be erroneously classified as menthol-insensitive cells (Fig. 9.4).

CTNs maintain an ongoing firing activity at normal skin temperature (33 °C), suggesting a tone of excitatory input of CTNs to the central cells where they synapse. This basal activity has been attributed to a temperature-dependent rhythmic oscillation of the membrane potential (Braun et al. 1980) (See also Chap. 10 by *Orio and Olivares* in this book). After the action potential firing frequency in response to a temperature drop reaches the peak (dynamic response), it slowly decreases to a lower sustained rate at the new temperature (static response) (Hensel and Iahn 1973) (see Fig. 10.3 in Chap. 10 of this book). The amplitude and rate of change of the cold-induced response is proportional to the rate and magnitude of the temperature decrease. The firing changes gradually from a beating to a bursting pattern at lower temperatures, and when the temperature is raised back, the electrical activity transiently silences with the rewarming process (Hensel and Iahn 1973; Darian-Smith et al. 1973).

In contrast to cold thermoreceptor neurons, C- and A $\delta$ -nociceptors responding to intense cold are silent at normal temperature of the skin, and fire only in response to temperatures below 15 °C. These neurons fire wide action potentials (>2 ms at the half-amplitude) with an inflection (or hump) in the falling phase, and also respond to other modalities of noxious stimuli (Bessou and Perl 1969; Croze et al. 1976;



**Fig. 9.4** Differential menthol sensitivity in low- and high-threshold cold thermoreceptor neurons. **a.** Simultaneous recordings of cold-induced  $[Ca^{2+}]_i$  and bath temperature in two cold-sensitive neurons with different temperature thresholds (low-threshold (LT-CS) and high-threshold (HT-CS)) during two consecutive cooling ramps, in control condition and in the presence of 100  $\mu$ M menthol. Note that in the low-threshold CSN menthol evoked a robust  $[Ca^{2+}]_i$  rise at 34 °C, while in the HT-CS neurons menthol only shifted the threshold of cold induced response to higher temperatures. **b.** Dot plot summarizing the effect of 100  $\mu$ M menthol on cold threshold in 86 trigeminal cold-sensitive neurons in culture. The neurons activated by 100  $\mu$ M menthol at 34 °C are marked by a magenta star. The menthol-insensitive neurons are marked by a green triangle. (Modified from Madrid et al. 2009)

LaMotte and Thalhammer 1982; Campero et al. 1996; Simone and Kajander 1996, 1997; Craig et al. 2001). Interestingly, additional nociceptors are excited with larger temperature reductions. Moreover, almost all primary somatosensory neurons respond to cold temperatures below 0°C, complicating the determination of the exact percentage of cold nociceptors in the peripheral somatosensory system (Simone and Kajander 1996, 1997; Campero et al. 2001).

In primary somatosensory neurons, TTx-insensitive Na<sub>v</sub>1.8 sodium channels are critical for noxious cold response. Unlike the TTx-sensitive Na<sup>+</sup> channels, whose slow inactivation is potentiated by cold, the inactivation of Na<sub>v</sub>1.8 is largely cold-insensitive. Thus, Na<sub>v</sub>1.8 channels allow the impulse generation of nociceptive neurons at very low temperatures (Zimmermann et al. 2007). In addition, the increase in membrane resistance and the decrease in the activation threshold of the Na<sup>+</sup> currents induced by cold augment the membrane potential change in response to a depolarizing stimulus.

## 9.3 TRPM8 Channels in Cold Transduction

### 9.3.1 *Molecular Cloning of TRPM8 and Expression in Primary Sensory Neurons*

**TRPM8** is the 8<sup>th</sup> member of the Melastatin-related family of **TRP** channels. The encoding gene was originally identified by its expression in normal prostate epithelial cells and prostate carcinomas, and the full-length cDNA of human TRPM8 was formerly called *trp-p8* (Tsavaler et al. 2001). The transcript corresponds to a 1104 amino acid protein with a marked homology to some members of the TRP channels superfamily. The expression of this protein in primary somatosensory neurons was not reported until 2002, when TRPM8 was cloned by two groups independently, and characterized as a cold- and menthol-activated ion channel (McKemy et al. 2002; Peier et al. 2002). In order to identify the putative cold and menthol receptor protein(s) responsible for cold and menthol sensitivity of CTNs, McKemy and co-workers constructed a cDNA expression library of trigeminal ganglia. They carried out a functional screening with this material, transfecting HEK293 cells with discrete cDNA pools obtained from this library. Using calcium imaging in isolated cells that exhibited intracellular calcium rises in response to menthol, they were able to identify a single cDNA sequence that conferred both menthol and cold sensitivity to these normally cold-insensitive cells, and classified it as a TRP channel (McKemy et al. 2002). Peier and colleagues, on the other hand, used a different strategy. Reasoning that the cold- and menthol-sensitive receptor could be a protein related to the recently cloned heat- and capsaicin-activated channel TRPV1 (Caterina et al. 1997), they searched directly for TRP-like proteins in genomic DNA databases looking for putative exons with similarity to the S4 and S6 transmembrane domains of TRPV1. One of these sequences was used to design primers to amplify a fragment of a puta-



tive TRP channel from a DRG cDNA library, and by using a rapid amplification of cDNA ends-PCR (RACE-PCR) in combination with an exon-prediction software, they obtained the full length sequence of the protein (Peier et al. 2002).

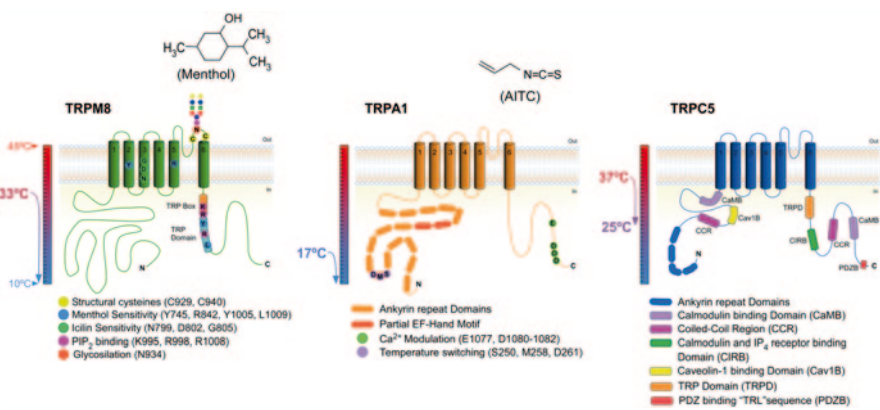
The TRPM8 channel is robustly expressed in trigeminal and dorsal root ganglia. TRPM8-positive neurons, corresponding to ~8–15% of the neurons of these sensory ganglia, exhibit a relatively diverse neural and biochemical phenotype. These neurons show immunoreactivity to peripherin, a marker of C-fibers, and intermediate filament NF200, a marker of A $\delta$  fibers. Furthermore, TRPM8 is co-expressed to a variable extent with nociceptive markers such as calcitonin gene-related peptide (CGRP), substance P, artemin receptor GFR $\alpha$ 3 and the nociceptive channel TRPV1 (Viana et al. 2002; Babes et al. 2004; Okazawa et al. 2004; Xing et al. 2006; Takashima et al. 2007; Dhaka et al. 2008; Axelsson et al. 2009; Parra et al. 2010; Zimmermann et al. 2011; Lippoldt et al. 2013). The NGF receptor TrkA is also expressed in a large subpopulation of TRPM8-positive neurons, and mediates the increase in cold-sensitivity of CTNs induced by this growth factor (Babes et al. 2004). A fraction of cold- and menthol-sensitive neurons also responds to proinflammatory mediators such as bradykinin, histamine and prostaglandin E<sub>2</sub>, with a reduction of its sensitivity to agonists (Linte et al. 2007; Zhang et al. 2012). On the other hand, non-peptidergic primary sensory neurons that bind isolectin B4 do not show significant expression levels of TRPM8 (Takashima et al. 2007). This neurochemical phenotype is in line with a sensory role of TRPM8 channels beyond innocuous cold thermosensation.

### 9.3.2 *Functional Properties and Modulation of TRPM8 Channels*

As mentioned before, TRPM8 is a nonselective cationic channel activated by cold, by natural and artificial cooling compounds such as menthol and icilin, and by voltage (for recent reviews, see (Babes et al. 2011; Latorre et al. 2011; McCoy et al. 2011; Yudin and Rohacs 2012; Almaraz et al. 2014; Madrid and Pertusa 2014)). Ion substitution experiments show a low discrimination of this channel among monovalent cations, but a significantly higher permeability for calcium ions ( $P_{Ca}/P_{Na}=3.2$ ;  $P_{K}/P_{Na}=1.1$ ;  $P_{Cs}/P_{K}=1.2$ ) (McKemy et al. 2002). Both cold and menthol increase the open probability of the channel. Activation of TRPM8 by cooling-mimetic compounds such as menthol explains the refreshing sensation evoked by natural and artificial activators of this channel, widely used in many toiletries. Interestingly, higher concentrations of menthol can evoke sensations of irritation and burning pain, probably due to the activation of TRPA1 channels in nociceptors (Karashima et al. 2007). TRPM8 shows a strong outward rectification at depolarized membrane potentials and an estimated single channel conductance of 60–90 pS depending on temperature. It is directly activated by cold, with a  $Q_{10}$  temperature coefficient as high as 25. Activation of the TRPM8 channel by low temperatures and by most of its natural and artificial agonists is related to a shift in its voltage-activation curve

to more negative voltages. This shift results in an increase in the open probability of the channel at physiologically relevant membrane potentials in response to thermal and chemical stimulation (Brauchi et al. 2004; Voets et al. 2004, 2007b).

TRPM8 is a homotetramer, where each subunit consists of six transmembrane domains with the N- and C- termini facing the cytosolic side of the plasma membrane (Fig. 9.5a). As in other members of the TRP channels family, the C-terminal domain of TRPM8 contains a TRP domain (Clapham 2003; Montell 2005). This domain plays an important role in its modulation by phosphatidylinositol 4,5-bisphosphate (PI(4,5)P<sub>2</sub>) (Rohacs et al. 2005) (see below). The coiled-coil motif located in the distal C-terminal domain plays an important role in the tetramerization and function of the channel (Erler et al. 2006; Tsuruda et al. 2006; Phelps and Gaudet 2007). Several regions have recently been described in the N-terminal domain as relevant for both cold and menthol sensitivity and proper folding and assembly of the channel protein (Pedretti et al. 2009; Pertusa et al. 2014). The pore module consists of the S5 and S6 domains and the interconnecting loop, where the selectivity filter is located. A unique N-glycosylation, at asparagine 934 facilitates the segregation of the channel to lipid rafts (Morenilla-Palao et al. 2009). This N-glycosylation site is flanked by two cysteine residues that form a conserved double cysteine motif (C929 and C940), which is essential for channel function (Dragoni et al. 2006). N-glycosylation of TRPM8 occurs in both native and recombinant channels, and it has a direct effect on their biophysical properties. The membrane potential for half-maximal activation ( $V_{1/2}$ ) of the channel carrying the single point mutation N934Q is 60 mV more positive than the wild type channel, shifting the mean temperature threshold of the cold response to lower temperatures (Pertusa et al. 2012) (see Chap. 3 by *Pertusa and Madrid* in this book).



**Fig. 9.5** Schematic representation of a TRPM8, TRPA1 and TRPC5 channel subunit. Left, TRPM8. Middle, TRPA1. Right, TRPC5. Each subunit presents six transmembrane domains, and the N- and C-terminal tails are located in the intracellular side of the plasma membrane. The most conspicuous functional domains and residues are indicated. Temperature-activation ranges and typical chemical activators are also depicted

Mean cold thresholds over 30°C can be found in corneal cold-sensitive nerve endings and cultured CTNs, as opposed to a mean value of 25°C observed in transfected hippocampal neurons or HEK293 cells heterologously expressing TRPM8 (Reid et al. 2002; Viana et al. 2002; de la Peña et al. 2005; Madrid et al. 2006, 2009; Malkia et al. 2007). At the same temperature, the  $V_{1/2}$  by equivalent cold stimulation is ~ 140 mV more negative in native TRPM8 than in recombinant channels. This results in a significantly lower thermal excitation threshold of CTNs compared to the heterologous expression systems such as HEK293 cells (Malkia et al. 2007). Thus, the activation of TRPM8 at physiological membrane potentials results in larger inward currents in cold-sensitive neurons than in recombinant systems. Interestingly, it has been reported different temperature activation threshold and menthol sensitivity among TRPM8 orthologs, which is coincident with the mean body temperature in endothermic animals and tuned to the environmental temperature of its ecological niche in ectotherms (Myers et al. 2009). Recently, Fujita and coworkers have also reported that warm ambient temperatures shift the cold threshold of TRPM8 to higher temperatures (Fujita et al. 2013).

Charge-neutralizing mutations of positively charged residues in the S4 transmembrane segment and the S4-S5 linker of TRPM8 cause a decrease in its voltage dependence, suggesting that this region may be part of the voltage sensor (Voets et al. 2007a). Nevertheless, despite the advances in the study of the voltage and temperature dependence of TRPM8, and in contrast to its activation by cooling compounds, both the precise location of the voltage sensor and the molecular determinants of the temperature sensitivity remain elusive.

Functional and biophysical properties of TRPM8 can be finely tuned by diverse mechanisms, including lipid-protein interactions, protein-protein interactions, activation of intracellular cascades of second messengers and post-translational modifications such as phosphorylation and N-glycosylation. These mechanisms can influence its contribution to cold sensing under physiological and physiopathological conditions (see (Almaraz et al. 2014; Madrid and Pertusa 2014) for recent reviews; see also Chap. 3 by *Pertusa and Madrid* in this Book). In fact, both cold and menthol responses of TRPM8 are reduced by its basal association with lipid rafts at the plasma membrane (Morenilla-Palao et al. 2009). Although still unexplored, these results suggest that the association of TRPM8 with specific membrane microdomains may affect the targeting and trafficking dynamics of this channel in the nerve endings of cold-sensitive neurons. On the other hand, PI(4,5)P<sub>2</sub> is a critical cofactor for TRPM8 activation, and the positions K995, R998 and R1008 within the TRP domain appear to mediate the interaction of this lipid with the channel (Rohacs et al. 2005). PI(4,5)P<sub>2</sub> is able to activate TRPM8 in the absence of other physical and chemical stimuli (Liu and Qin 2005; Rohacs et al. 2005). Similarly to cold and cooling compounds, positive modulation of TRPM8 by PI(4,5)P<sub>2</sub> seems to be related to a shift in the  $V_{1/2}$  value of voltage activation towards more negative membrane potentials (Daniels et al. 2009).

Desensitization of TRPM8 to sustained cold and menthol stimulation strongly depends on extracellular Ca<sup>2+</sup> (McKemy et al. 2002). It has been suggested that this property is related to a depletion of PI(4,5)P<sub>2</sub> of the plasma membrane. Cal-

cium entry through TRPM8 activates  $\text{Ca}^{2+}$ -dependent PLC, reducing the sensitivity of the channel to thermal and chemical stimulation by diminishing  $\text{PI}(4,5)\text{P}_2$  levels (Daniels et al. 2009). Sarria and coworkers on the other hand have proposed that tachyphylaxis in response to repetitive stimulation of TRPM8 would be mediated by PLC-dependent hydrolysis of  $\text{PI}(4,5)\text{P}_2$  and the activation of a PKC/Protein Phosphatase 1 cascade, and that the desensitization by single sustained stimulation of the channel would depend on calmodulin activation and  $\text{PI}(4,5)\text{P}_2$  availability (Sarria et al. 2011).

TRPM8 channel function can be also modulated by different protein kinases (see Chap. 3 by *Pertusa and Madrid* in this Book). These intracellular modulatory mechanisms could explain the desensitization of the channel induced by inflammatory mediators. Thus, it has been suggested that bradykinin reduces the cold and menthol responses of TRPM8 through the activation of Gq protein and PKC-dependent mechanisms, respectively (Premkumar et al. 2005; Linte et al. 2007). Nevertheless, recently Zhang and coworkers proposed an alternative explanation, suggesting that the inhibition of TRPM8 activity is due to a direct interaction of the  $\text{G}\alpha_q$  subunit with the channel, independent of the signaling pathway downstream of  $\text{G}\alpha_q$ -coupled receptors (Zhang et al. 2012).

### 9.3.3 *TRPM8 as the Main Molecular Entity in Cold Transduction*

In CTNs, the depolarizing inward cold-induced current ( $I_{\text{cold}}$ ) is the main responsible for the excitatory receptor potential in response to temperature reductions.  $I_{\text{cold}}$  displays a biophysical and pharmacological profile consistent with a depolarizing current depending on the TRPM8 channel. This current exhibits outward rectification and can be fully suppressed by BCTC, a potent blocker of TRPM8 (Madrid et al. 2006).

CTNs display a wide range of temperature thresholds. This differential thermal sensitivity is to a large extent determined by the functional counterbalance of two conductances with opposite effects on temperature-dependent excitability. Differential functional expression of TRPM8 (the main responsible for the excitatory cold-activated current) and  $\text{Kv}1.1$ - $1.2$  Shaker-like potassium channels (responsible for the excitability break current  $I_{\text{KD}}$ ) is intimately linked to the thermosensitive phenotype of individual CTNs. The fast-activating slow-inactivating outward  $I_{\text{KD}}$  current dampens the effect of the cold-induced depolarizing TRPM8-dependent current, shifting the temperature threshold of the neuron to higher values, reducing the net response of the CTN to temperature reductions. Interestingly, pharmacological suppression of  $I_{\text{KD}}$  induces cold-sensitivity in cold-insensitive neurons (Viana et al. 2002). It is important to mention here that a fraction of cold-sensitive neurons does not express TRPM8, suggesting the presence of other mechanisms. Closure of thermosensitive background  $\text{K}^+$  channels TREK-1 and TRAAK by temperature drops also contributes to increase the excitability of CTNs neurons under cold

stimulation (Reid and Flonta 2001; Viana et al. 2002; Noel et al. 2009). Recently, TASK-3 and Kv7.2-7.3 potassium channels have also been proposed as modulators of cold-sensitivity in TRPM8-expressing neurons (Vetter et al. 2013; Morenilla-Palao et al. 2014).

The use of genetically modified mice has been critical to unveil the role of TRPM8 in sensing innocuous and noxious cold. Mice lacking functional expression of TRPM8 were developed by three groups independently (Bautista et al. 2007; Colburn et al. 2007; Dhaka et al. 2007), and all three TRPM8 knockout strains presented a strongly impaired cold sensitivity. TRPM8 knockout mice failed to discriminate between cold and warm environments. These mice exhibited a clearly reduced avoidance to cold in two-temperature tests and in thermotaxis assays of temperature gradients, when compared to wild type animals (Bautista et al. 2007; Colburn et al. 2007; Dhaka et al. 2007). Molecular ablation of the TRPM8 channel also reduced the behavioral responses to cooling agents and abolished their response to systemic stimulation with icilin, one of the strongest chemical agonists of TRPM8 (Dhaka et al. 2007). Using calcium imaging techniques in cultured primary sensory neurons, Bautista and coworkers additionally found that TRPM8 knockout animals showed a strong decrease in the incidence and magnitude of cold-induced responses (Bautista et al. 2007). Extracellular recordings of single sensory fibers revealed that both low- and high-threshold cold-sensitive primary afferents have impaired responses to temperature reductions. Cold-sensitive C-fibers from TRPM8 knockout mice also display lower basal action potential firing, but no effect on their general excitability (Bautista et al. 2007). Accordingly, basal firing of corneal CTNs and their responses to cold and menthol are virtually absent in TRPM8 knockout mice and proportionally reduced in heterozygous animals (Parra et al. 2010). Thus, not only the static and dynamic responses to temperature drops, but also the ongoing spontaneous activity of CTNs is largely dependent of the functional expression level of the cold- and menthol-activated channel TRPM8.

More recently, Knowlton and coworkers demonstrated that selective ablation of TRPM8-expressing neurons yielded animals with an even higher cold insensitivity in both innocuous and noxious range than observed in TRPM8 knockout animals (Knowlton et al. 2013). This result not only supports a critical role of TRPM8-expressing neurons in cold sensing, but also suggests the participation of other molecular mechanisms in cold-detection. Animals that lack TRPM8-expressing neurons have normal mechanical and heat sensitivity, suggesting that these neurons could be largely dispensable for other somatosensory modalities (Knowlton et al. 2013). Using a similar ablation strategy to eliminate TRPM8-expressing neurons but in adult mice, Pogorzala and coworkers demonstrated that this approach strongly reduced the responses of these animals to cold in a wide range of low temperatures (Pogorzala et al. 2013).

Summarizing, molecular, cellular, biophysical and behavioral studies support a key role of TRPM8 in the molecular and neural machinery responsible for cold sensing in a wide range of low temperatures.

## 9.4 TRPA1 Channels in Cold Transduction

### 9.4.1 *Molecular Cloning of TRPA1 and Expression in Primary Sensory Neurons*

Transient Receptor Potential Ankyrin 1 channel (TRPA1), first identified in human fibroblasts and liposarcoma cells (Jaquemar et al. 1999), is the only member of the subfamily of TRPA channels found in mammals. A similar bioinformatic approach used in the molecular cloning of TRPM8 (Peier et al. 2002) was used by Patapoutian and his group to design primers for obtaining the full-length TRPA1 from mouse TG cDNA, encoding a 1115 amino acid protein (Story et al. 2003). TRPA1 was originally described as an ion channel activated by intense cold ( $< 17^{\circ}\text{C}$ ) (Story et al. 2003). TRPA1 works as a sensor of cell threat, responding to a plethora of structurally diverse pungent and noxious compounds, and nowadays it is seen as the main molecular entity in the somatosensory system responsible for the detection of irritant and potentially harmful substances. The TRPA1 channel also plays an important role as regulator of neuropeptide release and neurogenic inflammation, and is considered a promising molecular target for new analgesic and anti-inflammatory drugs (for reviews see (Bautista et al. 2006; Baraldi et al. 2010; Nilius et al. 2012; Zygmunt and Hogestatt 2014)). Moreover, TRPA1 functions as a detector of potentially toxic compounds throughout the animal kingdom, implying that it emerged early in evolution (Kang et al. 2010; Macpherson and Patapoutian 2010).

The proportion of sensory neurons expressing TRPA1 in sensory ganglia varies from one report to another, with observations ranging from 3.6 to 56.7% (Story et al. 2003; Jordt et al. 2004; Kobayashi et al. 2005; Nagata et al. 2005). TRPA1 can be found in both peptidergic (substance P- and CGRP-positive) and non-peptidergic (IB4-positive) primary sensory neurons (Hjerling-Leffler et al. 2007; Barabas et al. 2012). The channel colocalizes with the heat-activated polymodal channel TRPV1 in a subpopulation of nociceptors, while its co-expression with TRPM8 channels in cold thermoreceptor neurons is very low, comprising about 5% of TRPM8-expressing cells (Story et al. 2003; Jordt et al. 2004; Parra et al. 2010).

The particularly high threshold for the activation of TRPA1 by cold temperatures, along with its expression in nociceptive neurons, makes it a strong candidate as the molecular entity responsible for transducing painful cold.

### 9.4.2 *Functional Properties and Modulation of TRPA1 Channels*

TRPA1 is most likely a homotetramer, with each subunit having a topology of six transmembrane segments (S1-S6), a putative pore loop and selectivity filter between S5 and S6, and large cytoplasmic N- and C- terminal domains (Fig. 9.5b). TRPA1 is highly permeable to  $\text{Ca}^{2+}$ , with a PCa/PNa of 5.8 and 17% of the inward current carried by  $\text{Ca}^{2+}$  ions measured under physiological ionic conditions



(Karashima et al. 2010). TRPA1 is also activated by voltage, showing a shift in its voltage dependence in response to agonists toward more negative membrane potentials (Karashima et al. 2007; Zurborg et al. 2007). In cell-attached patches, TRPA1 shows conductances of  $\sim 65$  pS and  $\sim 110$  pS in the inward and outward directions respectively under physiological conditions. In the absence of divalent cations, TRPA1 shows a linear I-V relationship with a slope conductance of  $\sim 120$  pS (see (Nilius et al. 2011)). In heterologously expressed TRPA1 channels, cold produces an exponential increase in the open probability, which is accompanied by an important decrease in single channel conductance and an increase in the mean open time (Sawada et al. 2007; Karashima et al. 2009).

The molecular nature of TRPA1 agonists is remarkably diverse. The TRPA1 agonists include active ingredients in many natural compounds commonly used as spices in cooking: cinnamaldehyde present in cinnamon (Bandell et al. 2004), isothiocyanates in mustard oil, wasabi, and horseradish (Bandell et al. 2004; Jordt et al. 2004), allicin and diallyldisulfide in garlic (Bautista et al. 2005; Macpherson et al. 2005) and methyl salicylate in winter green oil (Bandell et al. 2004) are some examples. Noxious compounds like acrolein (in smoke and tear gas) (Bautista et al. 2006),  $\Delta 9$ -tetrahydrocannabinol (in marijuana) (Jordt et al. 2004), as well as irritant compounds generated endogenously during inflammatory, oxidative and nitritive stress responses, such as 4-hydroxynonenal, 15-deoxy- $\Delta 12,14$ -prostaglandin J2 and nitrooleic acid, have also been reported as activators of TRPA1 (Trevisani et al. 2007; Macpherson et al. 2007b; Taylor-Clark et al. 2008, 2009), just to name a few. Many agonists of TRPA1 are electrophilic compounds that activate the channel through covalent modification of cysteine or lysine residues (Hinman et al. 2006; Macpherson et al. 2007a; Takahashi et al. 2008; Bang and Hwang 2009). In contrast, non-electrophilic compounds that activate TRPA1 bind to the channel in a non-covalent manner (Karashima et al. 2007; Fajardo et al. 2008; Talavera et al. 2009; Nilius et al. 2012).

$\text{Ca}^{2+}$  ions modulate TRPA1 function. The N-terminal domain of this channel contains an EF-hand  $\text{Ca}^{2+}$  binding motif between Ankyrin Repeat Domain (ARD) 11 and 12, which has been related with  $\text{Ca}^{2+}$ -dependent modulation (Doerner et al. 2007; Zurborg et al. 2007).  $\text{Ca}^{2+}$  ions in the micromolar range induce a concentration-dependent leftward shift in the  $V_{1/2}$  toward physiologically relevant membrane potentials, enhancing the channels' chemical responses (Doerner et al. 2007; Zurborg et al. 2007). However, later studies suggest that apparently EF-hand motif is not required for this phenomenon (Wang et al. 2008).

Furthermore, it has been shown that activation and desensitization of TRPA1 channels in response to cold and chemical agonists are modulated by  $\text{Ca}^{2+}$  ions (Wang et al. 2008; Karashima et al. 2009). Thus, when  $\text{Ca}^{2+}$  is present, the current elicited by addition of the potent TRPA1 agonist mustard-oil shows a rapid activation and further desensitization. In the absence of  $\text{Ca}^{2+}$ , both processes are still present but with higher rise and decay time constants. An alternative  $\text{Ca}^{2+}$  binding site has been proposed within a cluster of acidic residues in the distal C-terminus of TRPA1, that has been found to be important for  $\text{Ca}^{2+}$ -mediated modulation of the agonist-induced responses (Sura et al. 2012).



### 9.4.3 TRPA1 as a Molecular Determinant in Noxious Cold Sensitivity

Cold-sensitivity of mammalian TRPA1 has been matter of intense debate. The seminal study of Story and co-workers described that mammalian cell lines and *Xenopus* oocytes expressing recombinant TRPA1 channels responded to intense cooling (Story et al. 2003). However, these results were not corroborated by similar experiments carried out by Jordt and co-workers one year later (Jordt et al. 2004). These authors studied thermal and chemical responses of cultured TG neurons, and they did not find a direct correlation between the cells responding to mustard oil and to noxious cold (Jordt et al. 2004). Later studies have argued either in favour (Bandell et al. 2004; Kwan et al. 2006; Sawada et al. 2007; Karashima et al. 2009) or against (Zurborg et al. 2007; Knowlton et al. 2010; Dunham et al. 2010) the cold sensitivity of TRPA1 and its contribution to cold sensing *in vivo*.

An alternative gating mechanism has been proposed, where the thermal sensitivity of TRPA1 would be an effect of cold-induced intracellular  $\text{Ca}^{2+}$ -increases required for channel activation in HEK293 cells (Zurborg et al. 2007; Caspani and Heppenstall 2009). Nevertheless, Karashima and co-workers found that heterologously expressed TRPA1 channels were able to respond to cold in a  $\text{Ca}^{2+}$ -independent fashion (Karashima et al. 2009). This cooling activation followed the same scheme of shifting  $V_{1/2}$  to more negative voltages, described as a general mechanism of thermal activation for other thermoTRP channels (Karashima et al. 2009).

It has been recently reported that differences in cold-sensitivity of TRPA1 channels, which appears to be largely species-dependent since mouse and rat TRPA1 can be activated by cold in heterologous expression systems but not its orthologous of monkey and human, would be related to differences in a single residue within the S5 domain. Moreover, this residue (G878 in rodent and V875 in primate) also appears underlie the different effects of menthol on the channel observed across species (Chen et al. 2013). Interestingly, Jabba and coworkers recently reported that three single point mutations (S250, M258 and D261) switch mouse TRPA1 channels from cold- to hot-sensitive without affecting its chemical sensitivity (Jabba et al. 2014).

The contribution of TRPA1 to cold-sensing *in vivo* has been studied using genetically modified animals. Thermal sensitivity of two different lines of transgenic TRPA1 null mice has been evaluated. Bautista and coworkers developed a TRPA1-deficient mouse through deletion of the putative pore-loop domain of the channel (Bautista et al. 2006). On the other hand, Kwan and coworkers, generated a TRPA1 null mouse deleting the exons downstream of the pore region, by insertion of an IRES cassette and polyadenylation sequences (Kwan et al. 2006). Although disruption of the TRPA1 gene abolishes the behavioral responses to chemical activators of the channel, it has no mayor effects in the response or prevalence of cold-sensitive neurons (Bautista et al. 2006; Kwan et al. 2006). Bautista and coworkers found that behavioral responses to noxious and innocuous cold stimuli between knockout and control mice were similar. In contrast, other studies using knockout mice show low-

er sensitivity to noxious and innocuous cold stimuli than wild type animals (Kwan et al. 2006; Gentry et al. 2010). Using the TRPA1 knockout mice developed by David Corey and his group, Karashima and coworkers reported a decreased cold sensitivity of null mutant mice, supporting the idea that TRPA1 plays a role in noxious cold detection *in vivo* (Karashima et al. 2009). A recent study on the TRPA1 knockout mice developed by Kwan and coworkers showed a considerable loss of nerve endings in the skin compared to wild type animals, which may account for the reduced responses to mechanical and cold stimuli of these mice *in vivo* (Andersson et al. 2013).

On the other hand, it has also been suggested that TRPA1 is a mediator of hypersensitivity to cold under pathological conditions (del Camino et al. 2010; Chen et al. 2011). It has also been reported that TRPA1 could play a role defining the thermal threshold in heat nociception (Hoffmann et al. 2013). In cultured vagal sensory neurons, Fajardo and coworkers showed a clear overlap between cold sensitivity of these neurons and TRPA1 pharmacology, and a large reduction in the population of cold-sensitive neurons from nodose ganglia in TRPA1 knockout mice compared with wild-type animals (Fajardo et al. 2008).

Thus, TRPA1 emerges as a critical component of the molecular machinery involved in the response to noxious agents in primary somatosensory neurons, with an important role in the release and detection of inflammatory mediators, and a species-dependent contribution to cutaneous thermal sensitivity under physiological conditions

## 9.5 TRPC5 Channel in Cold Transduction

**Transient Receptor Potential Canonical 5 (TRPC5)** (Fig. 9.5c) is a  $\text{Ca}^{2+}$ -permeable nonselective cation channel predominantly expressed in central nervous system, but is also found in primary somatosensory neurons and other cells and tissues, including vascular smooth muscle cells, endothelial cells, T-lymphocytes, monocytes cardiac cells, adrenal medulla, mammary glands and yolk sac, among others (reviewed by (Jiang et al. 2011; Zholos 2014)). This channel was cloned by Okada and colleagues in 1998 and formerly named TRP5 (Okada et al. 1998). Its full length mRNA sequence was obtained from the screening of a Oligo(dT)-primed cDNA library constructed with poly(A) RNA from adult mouse brain. This channel shows a wide diversity of opening mechanisms and activators. TRPC5 is mainly activated through receptors coupled to Gq, PLC and/or Gi proteins, but can be directly activated by a large list of agonist of the most diverse nature. This channel may also open in a store-dependent manner that requires other partner proteins, including other TRP channels (see (Zholos 2014)). It has been reported that TRPC5 also displays cold sensitivity in the mild to intense cold range (Zimmermann et al. 2011). Heterologous expression of TRPC5 in HEK293 cells generates cold-sensitive cation currents at temperatures below 37°C that can be potentiated by the PLC-Gq cascade activated by carbachol (Zimmermann et al. 2011). In addition to its sensitivity to cold, the expression pattern of this thermoTRP channel in primary somatosensory

neurons also pointed to a role in cold thermotransduction. Specifically, this channel is expressed in ~32% of mouse DRG neurons of small and medium diameter, in the nerve endings, axons and soma of these neurons, and in the regions I/II/III of superficial laminae of the dorsal horn of the spinal cord (Zimmermann et al. 2011).

A TRPC5 deficient mouse strain was developed by Riccio and co-workers in 2009, using a targeting construct that allows the deletion of the genomic region encoding amino acids 412–459 of the TRPC5 gene and the introduction of a frame shift and a stop codon (Riccio et al. 2009). Calcium imaging experiments using cultured DRG neurons from TRPC5 knockout mice revealed a significant reduction in the percentage of cold-sensitive neurons, with a decrease in both the menthol-sensitive and the menthol-insensitive populations of cold-responding neurons (Zimmermann et al. 2011). These results are in agreement with the reduction in the total population of TRPM8-positive neurons detected by immunostaining. However, in several behavioral tests, TRPC5 knockout mice have no impairment in their responses to cold in a wide range of temperatures, suggesting that TRPC5 is not required for innocuous and noxious cold detection (Zimmermann et al. 2011). The authors suggest that cold-sensitivity of TRPC5 could be involved in other adaptations, such as localized metabolic changes, vascular changes, retraction of neurites, or initiation of transcriptional programs, rather than cold thermotransduction.

## 9.6 Concluding Remarks

In summary, an increasing number of cellular, molecular, electrophysiological and behavioral studies support the idea that, among the thermoTRP channels activated by cold temperatures, TRPM8 is the main component of the molecular machinery responsible for cold sensitivity in primary somatosensory neurons. The uncontested involvement of TRPM8 in innocuous cold transduction is also being extended nowadays to noxious cold detection. TRPA1 appears to have a species-dependent contribution to cutaneous noxious-cold sensing under physiological conditions, and it is critical to the cold-sensitivity of visceral vagal afferents. On the other hand, further studies are necessary to determine if TRPC5 has a significant contribution to cold sensing. The molecular basis underlying cold-sensitivity of these thermoTRP channels is an exciting open question, and the whole picture of the role that these cold-activated thermoTRP channels play beyond thermosensation is still emerging.

**Acknowledgements** Supported by Grants CONICYT Anillo ACT-1113 (RM, MP, GU), FONDECYT 1131064 (RM), FONDECYT 11130144 (MP) and FONDECYT 3150431 (AG). We thank Dr. Carlos Belmonte and Dr. Annika Mäkiä for comments to the manuscript. We apologise for omission of relevant work due to space constraints. RP hold a PhD fellowship from CONICYT. We thank the support of VRIDEI-USACH.

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# Chapter 10

## Mathematical Modeling of TRPM8 and the Cold Thermoreceptors

Erick Olivares and Patricio Orio

**Abstract** The role of TRPM8 channel in thermotransduction involves several aspects of complexity that make it difficult to understand intuitively. First, it is activated by several stimuli (cold, voltage, agonists and intracellular signaling) that interact with each other, raising the question of how these interactions occur. Experimental evidence in this type of polymodal channel may be misinterpreted if the consequences of a working hypothesis are not considered carefully. Second, in parallel with the identification of TRPM8 as the main molecular transducer of cold temperatures in cold thermoreceptors of the somatosensory system, a list of other ion channels have been shown to be involved in the activity of cold-sensitive neurons and nerve endings. The variety of firing patterns observed at cold sensitive nerve endings arises from a complex interaction of ion channels that operate on different time scales. Mathematical modeling has been instrumental in understanding these phenomena, showing the consequences of the hypotheses raised. Here we review some of the models that have been proposed in these two areas: the activation of TRPM8 and TRPV1 by voltage and temperature, and the generation of firing patterns of cold thermoreceptors. We finish this chapter with a mathematical model showing how the calcium-dependent adaptation of TRPM8 may account for the response of cold thermoreceptors to rapid changes in temperature.

**Keywords** Mathematical modeling · Cold thermoreceptors · TRPM8

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## 10.1 Introduction

Mathematical modelling has been a useful tool for understanding neural behavior since the beginning of the neurosciences. Physically-tractable mathematical laws naturally arising from the electrical properties of cell membranes and their ion channels may be used to build models with a biological meaning. In turn, these are useful for interpreting experimental data and designing experiments. On the other hand, the physical laws that govern the behavior of ion channels are not completely clear but the mathematical representations are helpful for understanding them.

Typically, a mathematical model serves one or more purposes. The most common one is to test a hypothesis on the underlying mechanisms of an observable phenomenon. If the postulated mechanisms can be translated into a set of mathematical expressions, these expressions should reproduce the phenomenon; otherwise a different hypothesis will have to be considered. The model can make predictions that may be then tested experimentally, and in this way the underlying hypothesis is further challenged.

There are two issues related to the mathematical representation of the cold-activated channel TRPM8. First, its activation by temperature, voltage, and chemical agonists very early raised the question of whether these agents were acting on the same or different parts of the protein. The same question applies to other polymodal thermoTRP channels, among them, the related heat-sensitive TRPV1 channel. Various answers to the question were obtained, expressed by different mathematical representations and with predictions that could be assessed experimentally. Once the predictions were tested by the appropriate experiments, it was concluded that the different stimuli must be sensed by separate parts of the protein. In the second place, is the issue of the role of TRPM8 in the context of thermosensitivity. Cold-sensitive nerve endings and fibers have a complex behavior, firing action potentials in a variety of patterns depending on the temperature. Also, their activity is different depending on whether the temperature is raising or decreasing, being in fact better detectors of *temperature changes* than of *absolute temperature values*. Through the years, before and after the molecular cloning of TRPM8 was attained, many other ion channels have been involved in the function of cold thermoreceptor neurons. How do they all orchestrate to originate the complex responses of cold thermoreceptors? As we shall see in the second part of this chapter, mathematical modeling can help in the design of a hypothesis to be explored experimentally.

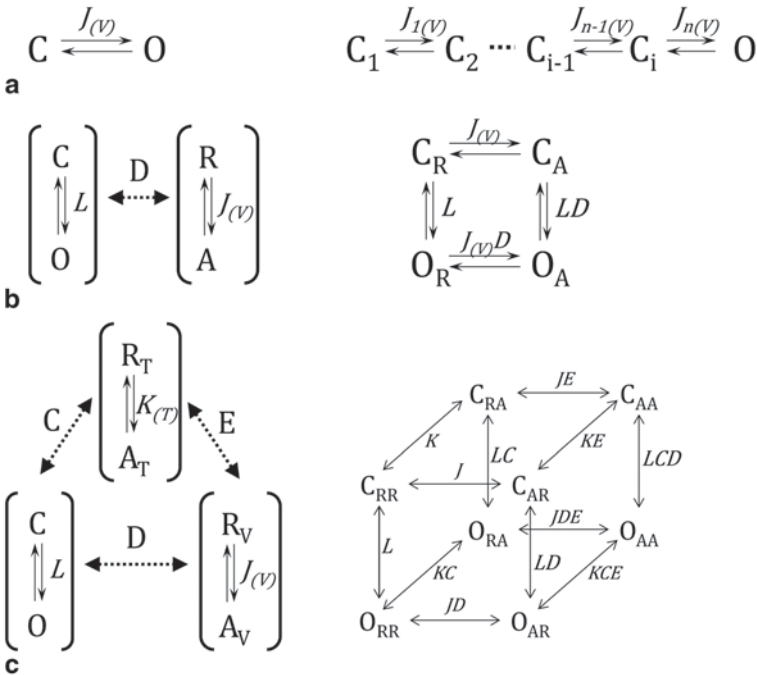
## 10.2 Allosteric Models and ThermoTRP Channel Activation

The activation of TRPM8 and TRPV1 channels raised the debate on the strictness of the coupling between the activators (temperature, membrane depolarization, etc.) and the pore gate. *Strict coupling* implies that whenever an activator is present the channel will open and this is the basic assumption for two-state models (Fig. 10.1a).

On the other hand, *allosteric coupling* (Fig. 10.1b, c) assumes that activation of the voltage (or other) sensor(s) does not lead directly to channel opening but rather to an increase of the open state probability. The difference may be subtle, but the two models have profound differences in their mechanistic interpretations and the prediction of channel behavior under certain conditions.

In a linear, strict coupling model, a maximum open probability of 1 is always attainable. This is seen from the open probability of a 2-state channel controlled by a voltage sensor:

$$P_O = \frac{O}{O+C} = \frac{1}{1 + \exp\left(-\frac{zF(V-V_0)}{RT}\right)}$$



**Fig. 10.1** **a** Simple two-state model. The voltage sensor and the channel gate are assumed to be the same molecular entity or two strictly coupled structures. Whenever the voltage sensor activates, the channels opens, and vice versa. **b** Allosteric activation by voltage. *Left*, two independent equilibria that interact allosterically. *R* and *A* represent the resting and activated state of the voltage sensor. *C* and *O* are the closed and open conformation of the pore gate. Only the equilibrium constant for voltage sensor activation, *J*, is voltage dependent. *Right*, when the combinations of the two equilibria are considered, a 4-state diagram results. *C<sub>n</sub>* and *O<sub>n</sub>* represent closed and open conformations of the pore, respectively, with the voltage sensors in the ‘n’ state. **c** Allosteric activation by voltage and temperature. A third equilibrium is added where *R<sub>T</sub>* and *A<sub>T</sub>* represent the resting and active conformations of the temperature sensor, respectively. Two new allosteric factors are introduced (*C* and *E*) to account for the possible interactions between structures. The expanded representation needs to be rendered in 3D (*right*)

In this expression and assuming  $z > 0$ , in the limit  $V \rightarrow +\infty$  we find that  $P_O \rightarrow 1$ . Conversely when  $V \rightarrow -\infty$ ,  $P_O \rightarrow 0$ . Similar expressions with the same consequences are found for schemes with many states, provided that the open state is only reached when all the sensors are activated (and conversely, the channel always opens when the sensors are active).

An allosteric gating scheme can be thought as two or more unconnected state transitions whose rate (and equilibrium) constants are modified depending on the state of the other equilibria. Fig. 10.1b depicts the simplest example of a two-state pore allosterically gated by a two-state voltage sensor. When the voltage sensor is in the resting state, ( $C_R$  or  $O_R$ ), the channel opens with a (probably very low) equilibrium constant  $L$ . When the sensor is in the activated state ( $C_A$  or  $O_A$ ), this equilibrium constant is multiplied by an allosteric factor  $D$ , thus incrementing the probability of the open state. Conversely, when the pore gate is in the open state, the equilibrium constant for the voltage sensor activation ( $J$ ) is multiplied by the same factor  $D$ , thus fulfilling the microscopic reversibility principle.

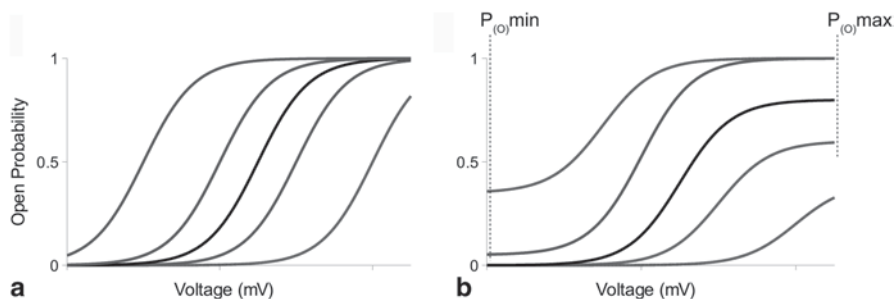
In the allosteric scheme depicted in (Fig. 10.1b), the open probability is

$$P_O = \frac{O_R + O_A}{C_R + C_A + O_R + O_A} = \left( 1 + \frac{1+J}{L(1+JD)} \right)^{-1}$$

In this case, if we assume that  $J$  increases with depolarization, in the limit  $V \rightarrow +\infty$  we find that  $J \gg 1 \Rightarrow P_O \rightarrow (1+LD^{-1})^{-1}$ . Conversely, when  $V \rightarrow -\infty$  we find that  $P_O \rightarrow (1+L^{-1})^{-1}$ , an expression that approaches  $L$  when  $L \ll 1$ . Thus, this scheme implies that minimum and maximum values for the open probability, different than 0 and 1, exist regardless of the state of the sensors. Fig. 10.2 graphically shows this difference in behavior of both strictly coupled and allosterically coupled models.

Other sensors for different stimuli are added to the model either as additional states or as another equilibrium that also interacts allosterically. Figure 10.1c shows the allosteric model originally proposed by Brauchi et al. (2004) for the activation by voltage and temperature, both in its allosteric representation and its expanded form that takes the shape of a cube. In this case a similar analysis of the minimum and maximum open probabilities results, where now the maximum open probability attainable by voltage stimulus depends not only on  $L$  and  $D$  but also on the temperature. At the same time, the maximum open probability that is reached by a temperature stimulus depends on the voltage.

In allosteric gated channels the number of states grows exponentially if other sensors such as for chemical agonists are included, and this type of scheme cannot be easily expressed in two dimensions. For instance, Matta and Ahern (2007) expanded the model to include activation by ligands. The expanded form requires four dimensions to be drawn with all the possible transitions. In the most recent and detailed study of TRPM8 activation by voltage and temperature, Raddatz et al. (2014) propose a three-tiered scheme with 75 states to account for all of their experimental observations (and this does not include activation by agonists). However, when seen in its allosteric depiction (as in Fig. 10.1c, left), it is evident that this type of activa-



**Fig. 10.2** Different behavior of strict coupling versus allosteric coupling. **a** A two-states model for voltage activation (strict coupling between sensor and gate) will always have maximum  $P_{(O)}=1$  and minimum  $P_{(O)}=0$ . Another agonist or activator can only change the  $V_{1/2}$  of the G/V relationship shifting the curve to the left or to the right. **b** On the other hand, with allosteric activation by voltage minimum and maximum  $P_{(O)}$  is not restricted to 0 and 1. In the model depicted in Fig. 10.1c, when the temperature sensor is resting ( $K(T)\ll 1$ ) the minimum  $P_{(O)}$  (when  $J(V)\ll 1$ ) is  $(1+1/L)^{-1}$  and the maximum open probability that can be reached by depolarization ( $J(V)\gg 1$ ) is  $(1+1/LD)^{-1}$ . On the contrary, when the temperature sensor is fully active ( $K(T)\gg 1$ ) the minimum  $P_{(O)}$  is  $(1+1/LC)^{-1}$  and the maximum is  $(1+1/LCD)^{-1}$ . The values of  $L$ ,  $C$  and  $D$  will determine whether these probabilities are significantly different from 0 or 1 for a macroscopic current analysis, but nevertheless they should always be measurable with single channel recordings

tion scheme has a relatively small number of parameter (compared to the number of states) and captures the proposed intramolecular interactions with simplicity.

The experimental behavior of a channel predicted by an allosteric gating scheme has several landmarks that have been described and extensively studied for the BK and HCN channels (Altomare et al. 2001; Horrigan and Aldrich 2002). They include the existence of several open states evidenced as multiple mean open times or current relaxations with multiple exponentials and the movement of gating charges between open states. However, as mentioned before, the most notable prediction are the minimum and maximum open probabilities that can be reached by hyperpolarization or depolarization. Depending on the values of  $L$  and  $D$  this may or may not be evident from macroscopic current recordings. For instance, the BK channel shows a maximum  $P_O$  almost equal to 1 ( $LD\gg 1$ ) and no minimum open probability because the equilibrium constant  $L$  has a weak voltage dependence that has been shown to be independent of the voltage sensors. Still, this phenomenon is evident at  $P_O < 10^{-6}$  (Horrigan et al. 1999).

(Brauchi et al. 2004) were the first to postulate an allosteric gating scheme for TRPM8, as their recordings showed that cold increases its maximum  $P_O$ . This was not noticed by (Voets et al. 2004) as they based their analysis and fitting only on macroscopic current recordings. On the other hand, Brauchi and coworkers measured actual  $P_O$  values with single channel recordings and non-stationary noise analysis. The observation was confirmed later for a wider range of temperatures and voltages, together with the finding of a *minimum*  $P_O$  at negative voltages which is also increased by cold (Raddatz et al. 2014). Matta and Ahern (2007) reported



the same for both TRPM8 and TRPV1 (in the case of TRPV1, heating increases maximum  $P_o$ ) also from single channel recordings. Moreover, Matta and Ahern showed that the minimum  $P_o$  of both channels is dramatically increased by the presence of chemical agonists (menthol for TRPM8, capsaicin and resiniferatoxin for TRPV1). The analysis presented by Voets et al. (2007) of the activation of TRPM8 by agonists led them to propose an allosteric gating scheme for the effect of menthol, while retaining a two-state mechanism for the activation by voltage (implying a strict coupling). The authors did not determine actual values of  $P_o$  and based their conclusions on normalized macroscopic G/V curves.

Not everything is about maximum and minimum open probabilities. Latorre and his group (Latorre et al. 2007; Raddatz et al. 2014) have shown that activation of TRPM8 by voltage, despite being very fast, shows a brief delay after the onset of a depolarizing pulse. Moreover, this delay is shortened by a depolarizing pre-pulse. This phenomenon, known as the Cole-Moore shift, cannot be observed in a two-states scheme for voltage activation and is a clear indication of multiple closed states (Cole and Moore 1960). More evidence supporting the existence of multiple open and closed states comes from single channel dwell time analysis in the TRPV1 channel (Liu et al. 2003), the bursting nature of the TRPM8 single channel activity (Zakharian et al. 2010), and the bi-exponential time course of TRPM8 current deactivation (Raddatz et al. 2014).

Finally, from a structural point of view, for a channel to be allosterically activated its sensors have to be different protein domains, also different from the pore gate. Though voltage and temperature sensors are far from being identified, a modular structure has been revealed in thermoTRP channels (Latorre et al. 2006; Brauchi et al. 2006), with growing evidence suggesting the existence of different activation domains for voltage, temperature and  $PIP_2$  in TRPM8.

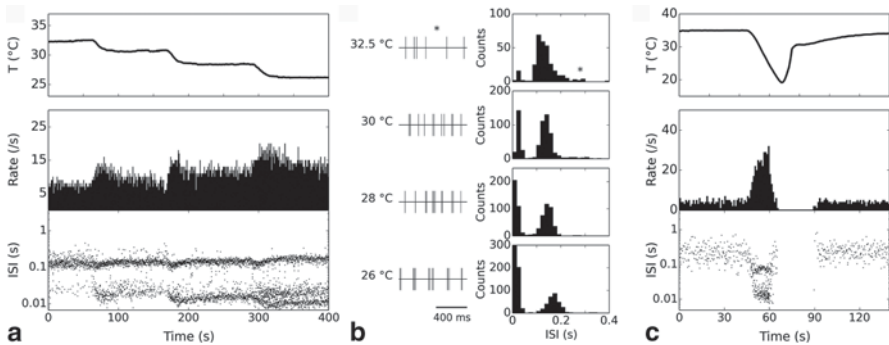
Thus, it appears difficult to sustain a strictly coupled two-state mechanism for voltage activation of thermo-TRP channels. A two-state model is simpler than allosteric models and reproduces satisfactorily the macroscopic conductance/voltage relationships and their modulation by temperature and agonists. However, the debate between strict and allosteric coupling of the voltage sensor and the pore gate has implications that go beyond the best fit of the experimental data. If the purpose of the model is to reproduce the channel behavior (for inserting it into a conductance-based neuron model, for instance) the choice will always be the simplest model. But when it comes to drawing mechanistic or structural conclusions, the model has to be challenged at every thinkable condition, especially at maximum and minimum  $P_o$ . Evidence for allosteric coupling has been found even in the family of Kv channels, largely thought to have the strictest coupling between the voltage sensor and the pore gate (Lu 2002). Still, allosteric coupling seems to be more convenient for channels activated by more than one stimulus, because the dynamic range for the effect of each agonist can be tuned independently.

### 10.3 TRPM8 Within the Big Picture of Cold Thermosensation

Since the cloning of TRPM8 (McKemy et al. 2002; Peier et al. 2002), this channel has been viewed as the main molecular sensor responsible for cold sensation. Its pivotal role was truly confirmed with the characterization of TRPM8 knock-out mice (Chung and Caterina 2007; Colburn et al. 2007; Dhaka et al. 2007; Bautista et al. 2007). However, more than 50 years of physiological characterization of cold thermoreceptors preceded the molecular findings (Hensel and Zotterman 1951; Iggo 1969; Hensel and Wurstler 1970; Dykes 1975; Bade et al. 1979). Based on electrophysiological and pharmacological evidence, other ion channels such as TREK-1/TRAAK have been proposed to play the role of transducing cold into electrical potentials, and growing molecular evidence shows that TRPM8 is a piece to be fit into a more complex picture.

#### 10.3.1 Cold Receptors as Pattern-Generating Machines

Mammalian thermoreceptor nerve endings that respond in the innocuous cold temperature range show a different behavior from other somatosensory receptors such as heat-thermoreceptors, mechanoreceptors or nociceptors. They are characterized by a regular ongoing spiking activity at normal skin temperature, that can be of tonic or bursting nature, and that is accelerated upon cooling the receptive field and suppressed by warming (Braun et al. 1980; Brock et al. 2001). The change of firing rate upon a temperature change suffers a strong adaptation in about a minute, relaxing to a steady state firing pattern that depends on the temperature (Fig. 10.3a, b). At low steady temperatures, the interval between spikes is increased but at the same time bursting is promoted so that single spiking events are rarely seen. On the contrary, at temperatures above 30–32°C single spikes prevail and ‘skipping’ events are evidenced as intervals that are twice the mean (or higher multiples) (Braun et al. 1980). It has been hypothesized that these regular firing patterns arise from a slow oscillation of the membrane voltage, on top of which action potentials are generated. Low temperature would prolong the period of this oscillation (by the typical effect of temperature on channel kinetics), thus allowing more action potentials per period at low temperatures (Braun et al. 1980). Current experimental evidence suggest that the slow oscillation is driven by a combination of slow TTX-resistant sodium channels (Brock et al. 2001) (probably Nav1.9 (Herzog et al. 2001)), and T-type voltage activated calcium channels (Schäfer et al. 1982). In addition, the HCN1 channel plays a role in setting the correct period of the slow oscillation. This channel is the main molecular counterpart of a hyperpolarization-activated current ( $I_h$ ) present in cold-sensitive neurons (Orio et al. 2009), and its blockade alters the firing pattern of cold-sensitive neurons (Orio et al. 2012).



**Fig. 10.3** Static and dynamic response of cold thermoreceptors. **a** Response of a corneal thermoreceptor ending to a step-like temperature protocol. Below the temperature trace, the firing rate and the Inter-Spike Intervals (*ISI*) are shown. *ISIs* are shown in logarithmic scale for a better depiction of bursts (low *ISI* values). **b** Representative firing patterns (*left*) and *ISI* histograms (*right*) from the data shown in **a**. Data for the histograms were taken after the transient responses has decayed. Asterisks denote skipping events, corresponding to intervals that are multiples of the main peak of the histogram. **c** Dynamic response of the thermoreceptor to an acute temperature pulse. Note the transient response of a bursting nature and how the nerve ending stops firing during the period of fast temperature raise. (Raw data obtained from Orio et al. 2012)

This variety of firing patterns is reproduced by mathematical models (Longtin and Hinzer 1996; Braun et al. 1998; Huber et al. 2000) based on a general model of slow wave bursting (Plant and Kim 1976). In this type of model, regular spiking or bursting is driven by a slow membrane potential oscillation and all that it takes to change the firing pattern with temperature decreases is to consider the usual effects of temperature on ion channels ( $Q_{10}$  values of 3 for kinetics and 1.3 for conductance). Huber and Braun's model was further modified to account for the experimental observations regarding  $I_h$  and HCN channels (Orio et al. 2012). In the latter work, the authors use mathematical modeling to support the hypothesis that the behavior of cold thermoreceptors from HCN1-knockout mice is best explained by compensation with the related channel HCN2 rather than by the absence of  $I_h$ . As a result, a new model of cold-sensitive nerve ending is proposed that not only reproduces the varying stationary firing patterns of cold thermoreceptors but also the role that HCN channels play in it. The model proposed by Orio and coworkers also introduces a TREK-1-like current, which is a background potassium channel that closes with decreasing temperature (Maingret et al. 2000; Kang et al. 2005). The decrease of a potassium conductance upon cooling was reported previously (Reid and Flonta 2001; Viana et al. 2002) and mice lacking both TREK1 and the similar TRAAK channels have altered cold perception (Noël et al. 2009).

Cold sensitivity in cold thermoreceptors is dampened by the expression of  $I_{KD}$ , an inhibitory outward slow-inactivating  $K^+$ -current (Viana et al. 2002; Madrid et al. 2009; see also Chap. 9 by González et al. in this book). The low activation threshold of  $I_{KD}$  and its slow inactivation implies that this current acts as an excitability brake that counteracts the depolarizing effect of cold in primary sensory neurons.

## 10.4 The Dynamic Response of Cold Thermoreceptors: A Model with TRPM8

The behavior of cold-sensitive nerve endings involves the participation of several ion channels, and the whole picture of cold transduction has yet to be unveiled. Where does TRPM8 fit in the cold transduction phenomenon? The mathematical models described before only reproduce the shifting patterns of nerve impulses that are observed when the cold receptors are already adapted to a steady temperature (the static response). They lack, however, the dynamic response, i.e. the acute changes in firing rate upon changes in temperature (Fig. 10.3c).

The TRPM8 channel is necessary for the response of cold thermoreceptors to acute changes in temperature. However, this does not explain all aspects of the dynamic response, such as the adaptation to a sustained stimulus or the absence of activity when heating. Yet the same ion channel may be at the center of all these phenomena. TRPM8 undergoes desensitization dependent on extracellular calcium (McKemy et al. 2002; Peier et al. 2002; Reid et al. 2002), by at least two mechanisms: one related to PIP<sub>2</sub> depletion by activated phospholipase C (Rohács et al. 2005; Daniels et al. 2009) and another one by Ca<sup>2+</sup>-Calmodulin which can revert the activation by PIP<sub>2</sub> (Sarria et al. 2011). The temporal scale of this process matches that of the adaptation of cold receptors, suggesting that TRPM8 desensitization can underlie the dynamic response in cold nerve endings. To test this hypothesis, we developed a model in which we inserted TRPM8 with a Ca<sup>2+</sup>-dependent adaptation into the existing Huber & Braun model.

As mentioned before, in the Huber & Braun model two slow currents (voltage activated sodium/calcium current and calcium activated potassium current) induce a slow subthreshold oscillation of the membrane potential. On top of this oscillation, a pair of fast depolarizing and repolarizing conductances originate action potentials whenever the membrane potential goes beyond the threshold. The kinetics of the channels, and thus the period of the oscillation, are modulated by temperature. This produces the variety of firing patterns observed in cold receptors. We inserted the TRPM8 conductance into the Huber & Braun model expecting it to maintain the variety of firing patterns, and additionally, that it could show the dynamic response of cold receptors.

Under these assumptions we have a modified Huber & Braun model in which the voltage membrane is given by

$$C_m \frac{dV}{dt} = -I_d - I_r - I_{sd} - I_{sr} - I_{M8} - I_{leak} + I_n$$

Where  $C_m$  is membrane capacitance;  $I_d$ ,  $I_r$ ,  $I_{sd}$  and  $I_{sr}$  are the depolarizing, repolarizing, slow depolarizing and slow repolarizing conductances, respectively;  $I_{M8}$  is the current mediated by the TRPM8 channel,  $I_{leak}$  corresponds to a unspecific leak current and  $I_n$  is a noise term. With the exception of  $I_{M8}$ , all the currents are present in the Huber & Braun model, and are described as follows:

$$I_i = \rho(T) g_i a_i (V - E_i) \quad i = d, r, sd, leak \quad \rho$$

$$I_{M8} = g_{M8} a_{M8} (V - E_i)$$

Where  $a_i \in [0,1]$  is an activation term,  $g_i$  is the maximal conductance parameter,  $E_i$  is the reversal potential for the current and  $\rho(T)$  is a temperature scaling factor for the currents.

The  $sr$  current is a calcium-activated channel, therefore its activation term actually reflects the internal calcium concentration. According to this, the current is expressed as

$$I_{sr} = \rho(T) g_{sr} \frac{a_{sr}^2}{a_{sr}^2 + 0.4^2} (V - E_{sr})$$

The activation terms  $a_r$  and  $a_{sd}$  follow the differential equation

$$\frac{da_i}{dt} = \phi(T) \frac{(a_i^\infty(V) - a_i)}{\tau_i} \quad a_i^\infty(V) = \frac{1}{1 + \exp(-s_i(V - V_{h,i}))} \quad i = r, sd$$

Where  $\phi(T)$  is a temperature factor for channel kinetics, and  $s_i$  and  $V_{h,i}$  are the parameters for the voltage-dependent activation of the currents.  $a_{sr}$  represents the internal calcium concentration, which increases with the inward  $I_{sd}$  current (a mixed sodium/calcium current) and decays according to the equation

$$\frac{da_{sr}}{dt} = \phi(T) \frac{-\eta I_{sd} - \kappa a_{sr}}{\tau_{sr}}$$

$a_d$  and  $a_{M8}$  have very fast kinetics compared to the other channels, therefore they are always equal to their steady-state values:

$$a_d = a_d^\infty(V) = \frac{1}{1 + \exp(-s_d(V - V_{h,d}))}$$

$$a_{M8} = a_{M8}^\infty(V, T) = \frac{1}{1 + \exp\left(-\frac{z_{M8}F}{RT}(V - V_h(T) - \delta V)\right)}$$

For the TRPM8 channel we have taken the simple two-state model proposed by Voets and colleagues (Voets et al. 2004; Voets et al. 2007) where  $z_{M8}$  is the voltage dependency and  $F, R, T$  have their usual physical constants meaning. The voltage for half-activation  $V_h$  is a function of temperature:

$$V_h(T) = \frac{CRT - (E_{a,C} - E_{a,O})}{z_{M8}F}$$

Where  $C$  is a constant related to the preexponential factors of the rate constants, and  $E_{a,C}$  and  $E_{a,O}$  are the activation energies for channel closing and opening, respectively. We introduced an additional shift  $\delta V$  that depends on intracellular calcium according to the equations

$$\frac{d\delta V}{dt} = \frac{\delta V^\infty (Ca^{2+}) - \delta V}{\tau_{\delta V}}$$

$$\delta V^\infty (Ca^{2+}) = \delta V_{min} + \frac{(\delta V_{max} - \delta V_{min}) [Ca^{2+}]}{[Ca^{2+}] + K_{Ca,M8}}$$

The intracellular calcium concentration is modeled considering that calcium influx to the cell is a fraction of  $I_{M8}$  and calcium removal follows a simple decay:

$$\frac{d[Ca^{2+}]}{dt} = -\frac{p_{Ca} I_{M8}}{2Fd} - \frac{[Ca^{2+}]}{\tau_{Ca}}$$

The temperature-dependent factors for conductance and kinetics are given, respectively, by:

$$\rho(T) = 1.3 \frac{T-25}{10}$$

$$\rho(T) = 3.0 \frac{T-25}{10}$$

Finally, the noise term was implemented as a low-pass filtered noise using an Ornstein-Uhlenbeck process with mean 0:

$$\frac{dI_n}{dt} = \frac{-I_n + D\xi(t)}{\tau_n}$$

Where  $\xi(t)$  is a normally distributed random variable (white noise) with zero mean and variance = 1.

Figure 10.4 shows the response of the model to an acute cold pulse using the following set of parameters:

$$g_d = 2.8 \text{ mS} / \text{cm}^2; g_r = 2.8 \text{ mS} / \text{cm}^2; g_{sd} = 0.2 \text{ mS} / \text{cm}^2;$$

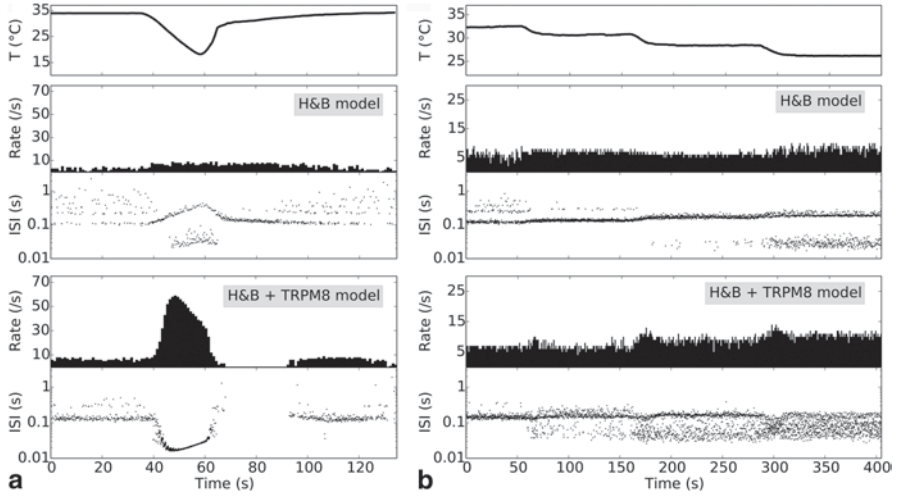
$$g_{sr} = 0.2 \text{ mS} / \text{cm}^2; g_{M8} = 2.74 \text{ mS} / \text{cm}^2; g_{leak} = 0.23 \text{ mS} / \text{cm}^2;$$

$$E_d = E_{sd} = 50 \text{ mV}; E_r = E_{sr} = -90 \text{ mV}; E_{leak} = -60 \text{ mV}; E_{M8} = 0 \text{ mV}$$

$$\tau_r = 2 \text{ ms}; \tau_{sd} = 10 \text{ ms}; \tau_{sr} = 35 \text{ ms};$$

$$V_{h,d} = V_{h,r} = -25 \text{ mV}; V_{h,sd} = -40 \text{ mV}; s_d = s_r = 0.25 \text{ mV}^{-1};$$

$$s_{sd} = 0.11 \text{ mV}^{(-1)}; \eta = 0.014 \text{ cm}^2 / \mu\text{A}; \kappa = 0.18,$$



**Fig. 10.4** Static and dynamic response of a thermoreceptor model including an adapting *TRPM8*. **a** Response of the original Huber & Braun (*H&B*) model (*middle*) and the *H&B* + *TRPM8* model (*bottom*) to an acute cold pulse (*top*; the same temperature trace as in Fig. 10.3c). Note that although the *H&B* model shows a slight increase in firing rate at lower temperatures, this response is instantaneous with respect to temperature and does not distinguish whether the temperature is decreasing or increasing. The *H&B* model shows a dramatic difference in firing rate depending on the direction of temperature change. **b** Response of the models to the temperature stimulus shown in (Fig. 10.3a). Although both models show a changing static firing pattern, the brief transient responses after each decrease in temperature are observed only in the *H&B* + *TRPM8* model

$$\begin{aligned}
 z_{M8} &= 0.82; C = 67.123; E_{a,C} = 173000J; E_{a,O} = 15700J; \\
 \tau_{\delta V} &= 18600 \text{ ms}; \delta V_{max} = 490 \text{ mV}; \delta V_{min} = -440 \text{ mV}; \\
 K_{Ca,M8} &= 25 \text{ nM}; p_{Ca} = 8.86 \times 10^{-6}; d = 1 \mu\text{m}; \tau_{Ca} = 27800 \text{ ms} \\
 \tau_n &= 1 \text{ ms}; D = \sqrt{0.0005 / dt}
 \end{aligned}$$

Equations were solved in the Neuron simulation environment (Hines and Carnevale 2001) with a fixed integration time step  $dt = 0.05 \text{ ms}$ .

It is important to note that we have chosen a two-state model to model *TRPM8*, in spite of the already discussed evidence in favor of multi-state models to reproduce its voltage- and temperature-dependence. The reason to do this is simple, and was already given at the end of the first section: the purpose of the model here is just to reproduce the behavior of the channels. A close look at some figures in the work of Raddatz and co-workers (Raddatz et al. 2014) reveals that *within the physiological values of voltage and temperature* (15–40 °C; -90–50 mV) the behavior of the channel is indistinguishable from a two-state model. Therefore, we chose the simplest model that fulfilled our purpose.



## 10.5 Concluding Remarks

Our model shows that the only element necessary to display a dynamic response in a model of cold thermoreceptor is a cold-activated channel (resembling TRPM8) with a slow activity-dependent adaptation. Moreover, and in tune with the discussion presented in the first part of this chapter, a simple two-state model of TRPM8 does the job, as all that is needed for this purpose is a phenomenological representation of the channel behavior within the physiological ranges of temperature and voltage.

**Acknowledgment** The work of the authors is supported by Fondecyt Grant 1130862, ACT-1113 and ACT-1104 (CONICYT, Chile) to PO. The Centro Interdisciplinario de Neurociencia de Valparaíso is a Millenium Science Institute supported by P09-022-F funds, Ministerio de Economía, Chile.

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# Index

## A

Abe, J., 84  
Abooj, M., 51  
Abrahamsen, B., 56  
Abraira, V.E., 148  
Abriel, H., 175  
Aebi, M., 76  
Agonists, 13, 16, 20, 55, 79, 81  
    of TRPA1, 198  
    of TRPM3, 174  
    of TRPV1, 43  
Ahern, G.P., 4, 15, 16, 20, 43, 81, 212, 213  
Akopian, A.N., 26  
Albert, J.T., 142  
Aldrich, R.W., 15, 17, 213  
Alessandri-Haber, N., 12, 50, 56, 155  
Al-Fageeh, M.B., 2  
Alloui, A., 177  
Almaraz, L., 78, 192  
Almeida, M.C., 167  
Altier, C., 84  
Altomare, C., 213  
Alvarez, D.F., 50  
Anand, P., 49  
Andersson, D.A., 22, 200  
Andrade, E.L., 56, 57  
Andre, E., 57  
Antagonists, 22, 44, 49, 55, 169  
Appendino, G., 43, 128  
Apweiler, R., 74  
Arendt, D., 98  
Arniges, M., 80  
Arshavsky, V.Y., 142  
Astorga, G., 102  
Avelino, A., 43  
Axel, R., 113, 117  
Axelsson, H.E., 192

## B

Babes, A., 5, 6, 54, 57, 192  
Bacigalupo, J., 104, 105, 114, 116  
Bade, H., 215  
Baez-Nieto, D., 4, 14, 20  
Bakalyar, H.A., 117  
Baker, H., 117, 118  
Bandell, M., 13, 22, 23, 56, 198, 199  
Bang, S., 47, 49, 198  
Banke, T.G., 20  
Barabas, M.E., 197  
Baraldi, P.G., 57, 58, 197  
Bargal, R., 10  
Barham, H.P., 121  
Baroiller, J.F., 2  
Barr-Gillespie, P.G., 147  
Barriere, D.A., 56  
Bartel, D.L., 128  
Bassi, M.T., 10  
Baum, M.J., 113, 116, 121  
Bautista, D., 18, 78  
Bautista, D.M., 13, 56, 151, 196, 197, 198,  
    199, 215  
Bavencoffe, A., 85  
Beck, A., 174  
Beech, D.J., 24, 51  
Bellringer, J.F., 116  
Belmonte, C., 187  
Bender, F.L., 171  
Benedikt, J., 20  
Benfenati, V., 173  
Berghard, A., 114  
Bessac, B.F., 56  
Bessou, P., 191  
Bevan, S., 45  
Bezzarides, V.J., 51  
Bharate, S.B., 55  
Bharate, S.S., 55

Bhave, G., 43, 81, 82, 83  
 Bidaux, G., 23, 54  
 Billig, G.M., 117  
 Birder, L., 50  
 Birnbaumer, L., 114  
 Bitter, 128, 134  
   perception of, 130  
 Blair, N.T., 24  
 Block, S.M., 142  
 Bloomquist, B.T., 103  
 Bodding, M., 13  
 Bohlen, C.J., 43  
 Boloskei, K., 154  
 Bonnington, J.K., 81  
 Boucher, Y., 136  
 Bourque, C.W., 154  
 Brauchi, S., 13, 14, 15, 16, 18, 21, 22, 23, 26,  
   193, 212, 213, 214  
 Braun, H.A., 190, 215, 216  
 Brayden, J.E., 155  
 Breaker, R.R., 4  
 Breer, H., 113, 117  
 Brennan, P.A., 113, 114  
 Brierley, S.M., 152  
 Broad, L.M., 43  
 Brock, J.A., 188, 215  
 Brunet, L.J., 118  
 Buck, L.B., 113, 117  
 Buck, L.B., 114  
 Burke, S.M., 121

## C

Calvo, R.R., 55  
 Campero, M., 191  
 Camprubi-Robles, M., 43, 46  
 Cao, E., 14, 17  
 Cao, X., 49  
 Carnevale, N.T., 220  
 Carstens, E., 136  
 Caspani, O., 199  
 Caterina, M.J., 5, 10, 12, 20, 43, 46, 76, 81,  
   148, 169, 170, 171, 191  
 Cesare, P., 10, 81, 170  
 Chai, S., 84  
 Chalfie, M., 142, 143, 144, 145  
 Chang, Q., 76  
 Chaudhari, N., 128  
 Chaudhari, S.S., 56  
 Chen, J., 13, 20, 26, 56, 58, 77, 199, 200  
 Chen, X.M., 168  
 Chianese, G., 58  
 Cho, H., 169, 177  
 Cho, Y., 55  
 Christensen, A.P., 142, 143, 151

Chuang, H.H., 13, 22, 55, 81  
 Chung, M.K., 5, 12, 14, 49, 77, 82, 171, 215  
 Chung, Y.D., 146  
 Chun, J.N., 27  
 Chyb, S., 103  
 Cinelli, A.R., 114  
 Ciura, S., 154  
 Ciurtin, C., 53  
 Clapham, D.E., 7, 8, 14, 22, 171, 172, 173,  
   193  
 Cohen, D.M., 76  
 Colburn, R.W., 13, 196, 215  
 Cold thermoreceptors, 187, 190, 215, 216  
   dynamic response, 217, 218, 220  
 Cold transduction, 79, 200, 201, 217  
   TRPA1 channels in, 197  
 Cole, K.S., 16, 214  
 Colledge, M., 84  
 Comelli, E.M., 76  
 Competitive, 44  
 Cordero-Morales, J.F., 14, 26  
 Corey, D.P., 114, 142, 143, 146, 147, 151  
 Cosens, D.J., 6, 101  
 Coste, B., 149, 169  
 Craig, A.D., 166, 191  
 Croze, S., 191  
 Cruz, A., 133  
 Csanady, L., 23  
 Cyert, M.S., 7

## D

Damak, S., 116, 130, 131, 134, 176  
 Daniels, R.L., 194, 195, 217  
 Darian-Smith, I., 190  
 Dattilo, M., 24  
 Davare, M.A., 51  
 Davis, J.B., 76, 81  
 D'Cotta, H., 2  
 De Araujo, I.E., 130  
 De Blas, G.A., 54  
 DeFalco, J., 58  
 Delany, N.S., 12  
 de la Peña, E., 194  
 Delay, R.J., 114  
 del Camino, D., 9, 56, 200  
 Delgado, R., 104, 105, 116  
 Denis, V., 7  
 De Petrocellis, L., 11, 49, 85  
 Devantier, H.R., 131  
 Devesa, I., 42  
 Dhaka, A., 10, 13, 14, 192, 196, 215  
 Dhennin-Duthille, I., 55  
 Dietrich, A., 76, 153  
 Di, M.V., 43

Ding, J., 51  
Distler, C., 84  
Docherty, R.J., 82, 83  
Doerner, J.F., 13, 47, 198  
Dorries, K.M., 113  
Dotson, C.D., 131  
Dragoni, I., 19, 20, 78, 193  
Drew, L.J., 148, 149  
Du, J., 174  
Dulac, C., 113  
Dunton, M.A., 50  
Dunham, J.P., 199  
Du, S., 152  
Dvoryanchikov, G., 128  
Dykes, R.W., 215

**E**

Earley, S., 49  
Eberl, D.F., 146  
Eberwine, J., 169  
Eddy, M.C., 131  
Eijkelkamp, N., 151  
Eisfeld, J., 12  
Enyedi, P., 176  
Erler, I., 23, 78, 193  
Erlund, I., 174  
Ernstrom, G.G., 142, 143  
Estacion, M., 104, 106  
Everaerts, W., 173

**F**

Facer, P., 47  
Fajardo, O., 23, 56, 198, 200  
Fan, H.C., 85, 86  
Fein, A., 105  
Felgines, C., 174  
Feng, P., 128  
Fernandes, J., 50  
Fernandez-Carvajal, A., 42  
Fernandez, J.A., 15, 17, 23  
Fernández-Peña, C., 55  
Ferrer, C., 99  
Ferrer-Montiel, A., 42, 49  
Finger, T.E., 113, 128  
Firestein, S., 113, 117  
Flemming, P.K., 51, 154  
Flonta, M.L., 196  
Flouris, A.D., 167  
Fonfria, E., 52  
Formaker, B.K., 134  
Frank, M.E., 134  
Franze, K., 106  
Frings, S., 117  
Fujita, F., 194

**G**

Galeotti, N., 49  
Galiotta, L.J., 177  
Gammie, S.C., 116  
Ganetzky, B., 9  
Gao, N., 131  
Gao, T., 84  
Gao, X., 50  
Garami, A., 171  
Garcia-Añoveros, J., 142, 146, 147  
Garcia-Martinez, C., 17, 45  
Garcia-Sanz, N., 21, 45  
Garger, A.V., 105  
Garrison, S.R., 153  
Gaudet, R., 9  
Gavva, N.R., 45, 171  
Gees, M., 129, 134, 136  
Geffeney, S.L., 146  
Gelstein, S., 121  
Gentry, C., 200  
Gesteland, R.C., 117  
Gillespie, P.G., 142, 143, 147  
Ginty, D.D., 148  
Gkika, D., 55  
Glendinning, J.I., 131  
Goel, M., 9  
Gold, G.H., 113, 117  
Goldstein, S.A., 176, 177  
Gomez, M.P., 105  
Gomis, A., 24, 51, 154  
Gong, Z., 9, 146  
Gonzalez-Silva, C., 117  
Goodman, M.B., 146  
Gottlieb, P., 153  
Gracheva, E.O., 26  
Grandl, J., 14, 16  
Grant, A.D., 155  
Green, B.G., 133  
Griffin, J.D., 169  
Grimm, C., 9, 12, 52, 53, 155, 174  
Grueber, W.B., 146  
Grunberg, N.E., 134  
Grycova, L., 22  
Gui, J., 58  
Guinamard, R., 13, 23, 175  
Guler, A.D., 50, 173  
Gutman, G.A., 20

**H**

Haga, J.H., 150  
Hahn, C., 150  
Halpern, M., 113  
Hamill, O.P., 142  
Hao, J., 148

Haraguchi, K., 174  
 Hara, Y., 173  
 Hardie, R.C., 7, 51, 101, 104, 106, 114, 116, 142  
 Hartneck, C., 53  
 Hasen, N.S., 116  
 Heat transduction, 4, 5  
 Heiner, I., 173  
 Helenius, A., 76  
 Hellwig, N., 19  
 He, L.P., 51  
 Hensel, H., 187, 188, 190, 215  
 Heppelmann, B., 187  
 Heppenstall, P.A., 199  
 Herzog, R.I., 215  
 Hicks, J.L., 101  
 Hille, B., 5  
 Hines, M.L., 220  
 Hinman, A., 198  
 Hinzer, K., 216  
 Hisatsune, C., 131  
 Hiura, A., 44  
 Hjerling-Leffler, J., 197  
 Hoffmann, T., 200  
 Hofmann, T., 9, 19, 106, 129, 131  
 Hogestatt, E.D., 197  
 Holakovska, B., 22  
 Holt, J.R., 147  
 Honore, E., 142  
 Honore, P., 154  
 Hori, A., 168  
 Hori, T., 2  
 Horrigan, F.T., 15, 17, 213  
 Hoshi, N., 84  
 Ho, T.C., 155  
 Hrabe de Angelis, M.H., 147  
 Huang, J., 43, 104, 105  
 Huang, S.M., 172, 173  
 Huang, Y.A., 128, 131  
 Huang, Y.J., 128, 131  
 Huber, A., 80, 86  
 Huber, M., 216  
 Huber, T.B., 154  
 Hudspeth, A.J., 142  
 Hu, G., 49  
 Hu, H.Z., 12, 49, 171, 172  
 Hu, J., 148  
 Hummel, T., 113, 121  
 Hurme, R., 2  
 Huynh, K.W., 21  
 Hwang, R.Y., 151  
 Hwang, S.W., 11

**I**

Iahn, M., 190  
 Iftinca, M., 169  
 Iggo, A., 215  
 Iida, T., 171  
 Ikeda, R., 149  
 Ikonen, E., 79  
 Inamura, K., 114  
 Innan, H., 114, 121  
 Inoue, K., 43  
 Inoue, R., 154  
 Iriuchijima, J., 187  
 Ishimaru, Y., 128  
 Islas, L.D., 4  
 Iwata, T., 114

**J**

Jabba, S., 199  
 Jahnel, R., 76, 77  
 Jang, Y., 175  
 Jaquemar, D., 197  
 Jara-Oseguera, A., 4  
 Jemal, I., 24  
 Jeske, N.A., 82, 83, 84  
 Jiang, L.H., 51, 200  
 Jin, M., 25, 50  
 Johnson, C.D., 54  
 Jones, D.T., 117  
 Jordt, S.E., 9, 13, 14, 20, 21, 56, 81, 82, 197, 198, 199  
 Julius, D., 18, 78, 81, 170  
 Jung, J., 82, 83  
 Jungnickel, M.K., 114  
 Jung, S., 51  
 Juvin, V., 46

**K**

Kajander, K.C., 191  
 Kang, D., 6, 169, 176, 216  
 Kang, K., 26, 197  
 Kang, N., 113  
 Kanosue, K., 168  
 Kapoor, M.D.S., 55  
 Karashima, Y., 5, 13, 20, 26, 56, 192, 198, 199, 200  
 Karlson, P., 113  
 Kaske, S., 116, 176  
 Katafuchi, T., 2  
 Katsumata, T., 128  
 Katz, B., 107, 142  
 Kawashima, Y., 147  
 Kedei, N., 76, 77  
 Keil, T.A., 146  
 Keller, A., 121



- Kelliher, K.R., 116  
 Kernan, M., 146, 156  
 Kerstein, P.C., 152  
 Keverne, E.B., 113, 114  
 Khairatkar-Joshi, N., 43  
 Kimchi, T., 116  
 Kim, J., 9, 146  
 Kim, K.X., 147  
 Kim, M., 216  
 Kim, M.T., 51  
 Kim, S., 114  
 Kim, S.E., 18, 147  
 Kindt, K.S., 151  
 Kinnamon, S.C., 128, 131, 142  
 Kiselyov, K., 10, 114  
 Kiyonaka, S., 51  
 Klausen, T.K., 50  
 Kleene, S.J., 117  
 Klose, C., 53  
 Knotkova, H., 44  
 Knowlton, W.M., 55, 196, 199  
 Kobayashi, K., 168, 170, 197  
 Kohler, R., 50  
 Koivisto, A., 58  
 Kolisek, M., 174  
 Koplak, P.A., 82  
 Kowase, T., 171  
 Kung, C., 143  
 Kurahashi, T., 117  
 Kwan, K.Y., 13, 26, 151, 152, 199, 200  
 Kym, P.R., 49
- L**
- Lamb, T.D., 98  
 LaMotte, R.H., 191  
 Lashinger, E.S.R., 55  
 Latorre, R., 4, 15, 16, 17, 18, 20, 26, 192, 214  
 Launay, P., 9, 19, 22  
 Lau, S.Y., 21  
 Lawson, J.J., 171  
 Lee, H., 173  
 Lee, N., 52, 174  
 Lee, Y.M., 51  
 Leffler, A., 46, 171  
 Lehen'kyi, V.Y., 55  
 Lehoux, S., 150  
 LeMasurier, M., 143  
 Lenzi, D., 103  
 Lesage, F., 177  
 Leung, H.T., 103  
 Levine, J.D., 56  
 Lev, S., 104  
 Lewin, G.R., 142, 148  
 Leybold, B.G., 115
- Liao, M., 18, 42  
 Liapi, A., 46  
 Liberles, S.D., 113, 117  
 Li, C., 150  
 Liedtke, W., 12, 50, 154, 155, 173  
 Light, 7, 51, 99, 101, 103  
 Li, L., 83  
 Liman, E.R., 112, 114, 116, 121, 128, 129,  
 131, 133, 176  
 Lingueglia, E., 149  
 Link, T.M., 171  
 Linte, R.M., 84, 85, 192, 195  
 Lin, W., 112, 113, 117, 118  
 Lipids, 9, 11, 12, 73  
 Lippoldt, E.K., 192  
 Lishko, P.V., 21  
 Littleton, J.T., 9  
 Liu, B., 13, 14, 17, 23, 55, 194, 214  
 Liu, D., 129, 134, 176  
 Liu, L., 134  
 Liu, P., 133  
 Liu, Y., 55  
 Li, W., 156  
 Longtin, A., 216  
 Lopez, F., 19, 118  
 Lopez, G.A., 17, 19  
 LopezJimenez, N.D., 128  
 Lorenzo, I.M., 173  
 Lou, D.-G., 99  
 Loukin, S.H., 50  
 Lowe, G., 113, 117  
 Lucas, P., 106, 114, 116  
 Luckhoff, A., 12  
 Ludeman, D.A., 25  
 Lumpkin, E.A., 148  
 Luscher, M., 113  
 Lu, Z., 214  
 Lyall, V., 128, 134
- M**
- Macpherson, L.J., 11, 13, 49, 197, 198  
 Madrid, R., 5, 6, 13, 56, 58, 78, 189, 192, 194,  
 195, 216  
 Maelkia, A., 55, 56  
 Maingret, F., 6, 169, 176, 177, 216  
 Majeed, Y., 53  
 Maksimovic, S., 148, 153  
 Malkia, A., 13, 22, 194  
 Ma, M., 113, 114, 117  
 Mandal, M., 4  
 Manning, A., 101  
 Manzoni, M., 10  
 Marcotti, W., 148  
 Maroto, R., 153

- Martinac, B., 142  
 Martinez, C., 166  
 Martinez-Marcos, A., 113  
 Masai, I., 105  
 Materazzi, S., 57  
 Mathematical modeling, 210, 216  
 Matsunami, H., 128  
 Matta, J.A., 4, 15, 16, 47, 212, 213  
 Matthews, J.M., 55  
 Ma, X.Y., 176  
 Ma, Y.Y., 173  
 McCarter, G.C., 148  
 McCoy, D.D., 78, 192  
 McDonnell, M.E., 45  
 McHugh, D., 23  
 McKemy, D.D., 5, 10, 13, 22, 23, 54, 55, 191, 192, 194, 215, 217  
 McNamara, F.N., 11  
 McNaughton, P.A., 10, 81, 82  
 Mechanotransduction, 10, 106, 169  
   mechanosensory, 150  
   models of, 144, 146, 147  
   principles of, 142, 143  
 Mechoulam, R., 49  
 Medhurst, A.D., 177  
 Meis, S., 51  
 Mei, Z.Z., 23  
 Menco, B.P., 117  
 Mendell, L.M., 81  
 Meotti, F.C., 58  
 Mercado, J., 22  
 Mercer, J.B., 2  
 Meseguer, V., 56  
 Meseguer, V.M., 133  
 Mihara, H., 46  
 Minke, B., 7, 51, 86, 101, 114  
 Miyamoto, T., 48  
 Mohapatra, D.P., 83  
 Monet, M., 46  
 Monteilh-Zoller, M.K., 19  
 Montell, C., 7, 9, 42, 45, 51, 101, 146, 170, 193  
 Moon, C., 142  
 Moore, J.W., 214  
 Moqrich, A., 12, 25, 49, 172  
 Morenilla-Palao, C., 43, 78, 79, 82, 193, 194, 196  
 Morgan, C.P., 147  
 Morita, H., 155  
 Morris, C.E., 142  
 Morrison, S.F., 167  
 Moshourab, R., 142  
 Moussaieff, A., 49  
 Mueller, K.L., 128  
 Muller, U., 147  
 Munger, S.D., 113, 114, 117  
 Muñoz, Y., 103  
 Muraki, K., 12, 46, 154, 171  
 Myers, B.R., 18, 194  
  
**N**  
 Nadler, M.J., 9  
 Nagamine, K., 173  
 Nagata, K., 151, 197  
 Nakamura, K., 168  
 Nakamura, T., 113, 117  
 Nakayama, T., 168  
 Nara, T., 5  
 Nasi, E., 105  
 Nath, A.K., 51  
 Nau, C., 83  
 Nauli, S.M., 10  
 Naylor, J., 53, 174  
 Naziroglu, M., 174  
 Neeper, M.P., 46, 171  
 Nelson, D.L., 3, 4  
 N-glycosylation, 74, 194  
   modulation of, 74, 76  
   TRPM8, 78, 79  
   TRPV1, 76, 77  
   TRPV4, 79, 80  
 Nilius, B., 7, 14, 17, 19, 23, 24, 25, 46, 50, 80, 128, 133, 142, 143, 151, 175, 197, 198  
 Noel, J., 177, 196, 216  
 Nolan, P.M., 147  
 Non-competitive, 44, 45  
 Nowycky, M.C., 20  
 Numata, T., 155  
 Numazaki, M., 21, 81  
  
**O**  
 Oberwinkler, J., 12, 52, 174  
 Obukhov, A.G., 20  
 Ogura, T., 113, 116  
 O'Hagan, R., 142  
 Öhrwall, H., 134  
 Oike, H., 134  
 Okada, T., 51, 106, 200  
 Okamoto, Y., 55  
 Oka, Y., 131  
 Okazawa, M., 192  
 Okubo, K., 152  
 Olfaction, 127, 142  
 Oliveira-Maia, A.J., 136  
 Oliveria, S.F., 84  
 Ordaz, B., 24

- Ordinary odorants, 113  
 Orio, P., 6, 17, 189, 215  
 Orr, A.W., 150  
 Ortar, G., 58  
 Ouadid-Ahidouch, H., 55  
 Owsianik, G., 7, 17, 19, 25, 46, 170, 171, 172, 174
- P**
- Pain, 7, 43, 51, 55, 58, 76, 87, 147, 187  
 Palmer, C.P., 7  
 Palmer, R.K., 175  
 Pan, B., 147  
 Panin, V.M., 76  
 Pan, Z., 173  
 Papakosta, M., 18  
 Pareek, T.K., 83  
 Parks, D.J., 55  
 Park, U., 12, 46, 154, 172  
 Parnas, M., 86, 106, 114  
 Parra, A., 55, 192, 196, 197  
 Pathology, 42, 43, 55, 149  
 Paul, M., 49  
 Payne, R., 105  
 Pedemonte, N., 177  
 Pedersen, S.F., 143  
 Pedretti, A., 193  
 Peier, A.M., 5, 10, 12, 13, 47, 54, 172, 191, 192, 197, 215, 217  
 Peng, H., 85  
 Penna, A., 46  
 Peralvarez-Marin, A., 46  
 Pereira, V., 177  
 Perez, C.A., 129, 176  
 Perl, E.R., 191  
 Perozo, E.I., 106  
 Perraud, A.L., 9, 19, 22, 173  
 Pertusa, M., 20, 78, 79, 189, 192, 193, 194  
 Peterlin, Z., 56  
 Petersen, C.C., 7  
 Petrus, M., 152  
 Phelps, C.B., 21, 22, 193  
 Pheromones, 7, 112, 121  
 Philippaert, K.P., 136  
 Phillipp, S.E., 12  
 Phillips, A.M., 6  
 Phosphorylation, 21, 23, 24, 74, 80, 84, 85, 87, 143  
   of TRPV1, 82, 83, 84  
   role of, 81  
 Photoreceptor, 98, 100, 101, 103  
 Pieau, C., 2  
 Planells-Cases, R., 44  
 Plant, R.E., 216  
 Plant, T.D., 51  
 Pochynyuk, O., 50  
 Pogorzala, L.A., 171, 196  
 Pollema-Mays, S.L., 177  
 Ponsati, B., 45  
 Popescu, D.C., 86  
 Postma, M., 142  
 Post-translational modifications (PTMs), 73, 87, 194  
 Prawitt, D., 129, 131, 175  
 Premkumar, L.S., 17, 51, 81, 84, 195  
 Prescott, E.D., 81  
 Preti, G., 121  
 Prevorskaya, N., 55  
 Primary sensory neurons, 54, 186  
   expression in, 191, 192, 197  
 Pristera, A., 79  
 Prober, D.A., 26, 151  
 Prole, D.L., 25  
 Pro-Sistiaga, P., 113  
 Provencio, I., 99  
 Pulido, C., 99  
 Pumpllin, D.W., 128
- Q**
- Qin, F., 194  
 Qin, N., 12, 46, 55, 171  
 Quick, K., 153
- R**
- Raddatz, N., 212, 213, 214, 220  
 Ramachandran, R., 55  
 Ramsey, I.S., 7, 21  
 Ranade, S.S., 150  
 Rao, V.S., 49  
 Rasche, S., 117  
 Rathee, P.K., 84  
 Rathner, J.A., 168  
 Reading, S.A., 155  
 Reed, R.R., 117, 118  
 Reid, G., 189, 194, 196, 216, 217  
 Reilly, R.M., 49  
 Ren, Z., 131  
 Restrepo, D., 113, 114, 117  
 Retina, 98, 105  
 Reuss, H., 104, 107  
 Rhen, M., 2  
 Riccio, A., 51, 201  
 Riera, C.E., 132  
 Rohacs, T., 13, 23, 192, 193, 194, 217  
 Romanov, R.A., 128, 130  
 Romanovsky, A., 169  
 Ronnett, G.V., 142

Roper, S.D., 128, 134  
 Rosenbaum, T., 21  
 Ross, R.A., 11  
 Roza, C., 149  
 Rubin, G.M., 101  
 Runnels, L.W., 9  
 Runnenburger, K., 114  
 Ryu, S., 19, 20

## S

Sabnis, A.S., 54  
 Sachs, F., 142  
 Saito, S., 25, 26  
 Salas, M.M., 26  
 Salazar, H., 11  
 Salt, 128, 131  
 Samad, A., 24  
 Samer, R.E., 58  
 Sandoz, G., 84  
 Sano, Y., 9  
 Santos, F.A., 49  
 Sardar, P., 25  
 Sarria, I., 195, 217  
 Sawada, Y., 198, 199  
 Saxton, T.K., 121  
 Schachter, H., 76  
 Schaefer, M., 9, 51  
 Schäfer, K., 215  
 Schild, D., 113, 114, 117  
 Schlingmann, K.P., 9  
 Schnitzler, M., 154  
 Schnizler, K., 84  
 Schoppa, N.E., 14  
 Schultz, G., 53  
 Schwartz, M.A., 150  
 Scott, H., 76  
 Scott, K., 101, 102  
 Semiochemical, 117, 120  
   definition of, 113  
 Semtner, M., 24  
 Sensory systems, 112  
 Sensory transduction, 4  
 Sharif-Naeini, R., 154  
 Shearer, A.E., 147  
 Shepherd, G.M., 142  
 Sherkheli, M.A., 47, 49  
 Shigetomi, E., 58  
 Shimizu, S., 24, 51  
 Shingai, R., 25  
 Shin, J.B., 156  
 Shu, X., 81  
 Shyu, K.G., 150  
 Sidi, S., 9, 156  
 Sigworth, F.J., 14

Simone, D.A., 191  
 Simons, C.T., 134, 136  
 Simons, K., 79  
 Smales, C.M., 2  
 Smith, F.D., 84  
 Smith, G.D., 5, 12, 47, 172  
 Smith, P.L., 50, 173  
 Smith, R.J., 147  
 Son, A.R., 155  
 Sour, 128  
 Soya, M., 152  
 Spassova, M.A., 153  
 Spehr, M., 114, 116  
 Starkus, J.G., 174  
 Staruschenko, A., 26  
 Stayner, C., 10  
 Stephan, A.B., 117  
 Stewart, A.P., 21  
 Stohr, H., 117  
 Story, G.M., 5, 9, 13, 20, 56, 197, 199  
 Stotz, S.C., 49  
 Stowers, L., 115  
 Straub, I., 53, 174  
 Streng, T., 152  
 Strotmann, R., 12, 21, 50, 154, 173  
 Stucky, C.L., 152  
 Studer, M., 81, 82  
 Sukharev, S., 142  
 Sukumar, P., 51  
 Sung, T.S., 24  
 Sun, M., 10  
 Sura, L., 24, 198  
 Susankova, K., 18  
 Suzuki, M., 155  
 Sweet, 120, 128, 130, 131, 134, 176  
 Swope, S.L., 80  
 Szallasi, A., 43, 44  
 Szelenyi, Z., 171

## T

Tabuchi, K., 50, 155  
 Tajino, K., 167  
 Takahashi, N., 198  
 Takaiishi, M., 58  
 Takashima, Y., 192  
 Takezawa, R., 175  
 Talavera, K., 5, 10, 120, 129, 130, 133, 134,  
   136, 175, 176, 198  
 Tamayo, N., 171  
 Tamayo, N.A., 55  
 Taniguchi, M., 114  
 Taruno, A., 128, 131  
 Taste transduction, 116, 128, 130, 133, 136  
 Taylor-Clark, T.E., 57, 198

- Taylor, C.W., 25
- Teicher, M.H., 113
- Thalhammer, J.G., 191
- Therapeutic index, 44
- Thermoregulation, 165, 166, 173, 178  
  sensory afferent input, 167
- Thermosensation, 5, 25, 80, 166, 192, 215
- Thermo TRP channels, 10, 15, 17  
  models of, 20, 21, 22
- Thiel, G., 53, 174
- Thompson, J.A., 113
- Thomson, J.N., 144
- Thorneloe, K.S., 50
- Tizzano, M., 113
- Todaka, H., 12
- Todd, A.J., 166
- Togashi, K., 5, 10, 173, 174
- Tomassini, S., 134
- Tominaga, M., 11, 43, 134, 170
- Tong, Q., 23
- Topala, C.N., 19
- Tordoff, M.G., 132
- Torello, A.T., 113
- Torocsik, B., 23
- Torre, V., 142
- Touch, 7, 142, 145, 148, 155, 187
- Transduction, 4, 7, 27, 74, 99, 102, 113, 118,  
  120, 133, 134, 200
- Transient receptor potential ankyrin 1  
  (TRPA1), 5, 9, 14, 20, 42, 56, 58,  
  134, 151, 152, 199  
  modulation of, 197, 198
- Transient receptor potential canonical 5  
  (TRPC5), 5, 13, 14, 20, 24, 51, 154,  
  200
- Transient receptor potential channel M5  
  (TRPM5), 112, 113, 116, 117, 129  
  effects of, 118, 119  
  role of, 130
- Transient receptor potential channel M8  
  (TRPM8), 5, 10, 13, 14, 16, 18, 22,  
  49, 54, 55, 78, 79, 191  
  modulation of, 192, 193, 194, 195
- Transient receptor potential (TRP) channel, 5,  
  14, 17, 20, 103, 113  
  allosteric-polymodal, 15  
  history of, 6  
  mechanosensory, 150  
  modulation of, 74  
  pharmacology of, 42  
  physiology, 27  
  regulation of, 80, 81  
  superfamily of, 7  
  thermosensitive, evolution of, 25, 26
- Transient receptor potential vanilloid 1  
  (TRPV1), 5, 9, 10, 15, 19, 43, 84,  
  154, 170, 171  
  basal phosphorylation of, 82, 83
- Trebak, M., 106
- Treesukosol, Y., 128
- Trevisani, M., 11, 198
- Trotier, D., 121
- TRP channels  
  pharmacology of, 42
- TRPP3/PKD1L3, 127, 128
- Tsavalier, L., 54, 55, 191
- Tse-Dinh, Y.C., 4
- Tsunoda, S., 101
- Tsuruda, P.R., 23, 78, 193
- U**
- Uchida, K., 134, 174
- Ueda, K., 43
- Ufret-Vincenty, C.A., 22
- Ullrich, N.D., 129, 175
- Umami, 128, 130, 136
- Umezū, T., 49
- Uncompetitive, 44, 45
- V**
- Valdes-Rodriguez, R., 46
- Valente, P., 45
- Valero, M.L., 55
- Vallin, K.S.A., 58
- Van Buren, J.J., 43
- Vandenbeuch, A., 128, 131
- Van Der Stelt, M., 43
- Vandewauw, I., 175, 176
- Vannier, B., 114
- Varga, A., 43
- Varki, A., 74, 75
- Vay, L., 169
- Veldhuis, N.A., 76, 78
- Vellani, V., 43, 81
- Venkatachalam, K., 10, 14, 45, 146, 170
- Vennekens, R., 49
- Vetter, I., 196
- Viana, F., 55, 189, 192, 194, 195, 216
- Vidal-Mosquera, M., 45
- Vigh, L., 4
- Vilceanu, D., 152
- Vincent, F., 50
- Vision, 7, 87, 98, 99
- Viswanath, V., 26
- Vlachova, V., 14, 21, 22, 170
- Voets, T., 11, 13, 14, 15, 17, 22, 129, 133, 143,  
  169, 170, 193, 194, 213, 214, 218

Vogt-Eisele, A.K., 49  
 Vomeronasal, 112, 113, 114, 120  
 Woolstra, O., 80, 87  
 Vriens, J., 5, 9, 46, 49, 50, 53, 85, 155, 174,  
 175

**W**

Wagner, T.F., 12, 53, 174  
 Wahl, P., 45  
 Walder, R.Y., 5, 9  
 Waldmann, R., 169  
 Walker, K.M., 45  
 Walker, R.G., 9, 142, 146, 156  
 Wang, G., 26  
 Wang, H., 20  
 Wang, X., 43  
 Wang, Y.Y., 55, 198  
 Watanabe, H., 5, 12, 23, 50  
 Wechselberger, M., 169  
 Wegierski, T., 85, 86  
 Wehage, E., 173  
 Wei, H., 56, 152  
 Weinbaum, S., 150  
 Wes, P.D., 7, 8, 51, 101  
 Willette, R.N., 173  
 Wirkner, K., 19, 21, 77  
 Wissenbach, U., 12, 154  
 Wittbrodt, J., 98  
 Witt, M., 113, 121  
 Wolstenholme, A.J., 25  
 Wong, C.O., 51  
 Woodbury, C.J., 171  
 Wood, J.N., 46  
 Woo, S.H., 149  
 Wuensch, T., 51  
 Wu, L.J., 51  
 Wurster, R.D., 215  
 Wyart, C., 121  
 Wysocki, C.J., 121

**X**

Xiao, B., 169  
 Xiao, R., 21  
 Xing, H., 55, 192  
 Xu, C., 173  
 Xue, T., 99  
 Xu, H., 5, 11, 12, 47, 49, 80, 85, 172  
 Xu, Q., 150  
 Xu, S.Z., 51, 53

**Y**

Yamamura, H., 55  
 Yang, C., 114  
 Yang, F., 15, 17, 18  
 Yang, X.R., 50  
 Yang, Y.D., 177  
 Yan, Z., 147  
 Yao, J., 14  
 Yau, K.W., 117  
 Yee, N.S., 55  
 Yeh, B.I., 19  
 Yildirim, E., 114  
 Yoon, I.S., 173  
 Yoshida, T., 24  
 Yudin, Y., 192

**Z**

Zakharian, E., 14, 214  
 Zamudio-Bulcock, P.A., 53  
 Zeng, F., 24, 51  
 Zhang, D., 171  
 Zhang, F., 21  
 Zhang, J.J., 114  
 Zhang, P., 114, 116, 120  
 Zhang, W., 147  
 Zhang, X., 56, 76, 81, 82, 84, 117, 192, 195  
 Zhang, X.F., 152  
 Zhang, Y., 116, 130, 131  
 Zhang, Y.H., 168  
 Zhang, Z., 129, 174  
 Zhao, H., 118  
 Zholos, A.V., 200  
 Zhou, J., 10  
 Zhu, B., 55  
 Zhu, X., 7, 8  
 Zimmermann, K., 5, 10, 13, 20, 24, 191, 192,  
 200, 201  
 Zotterman, Y., 188, 190, 215  
 Zufall, F., 51, 114, 116  
 Zuker, C., 146  
 Zuker, C.S., 86, 101, 102  
 Zurborg, S., 13, 198, 199  
 Zygmunt, P.M., 197