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Trieste, Italy



Functional characterization of molecular candidates for the calcium-activated chloride channels in the cilia of olfactory sensory neurons.

Thesis submitted for the degree of "Doctor Philosophiae"

SUPERVISOR

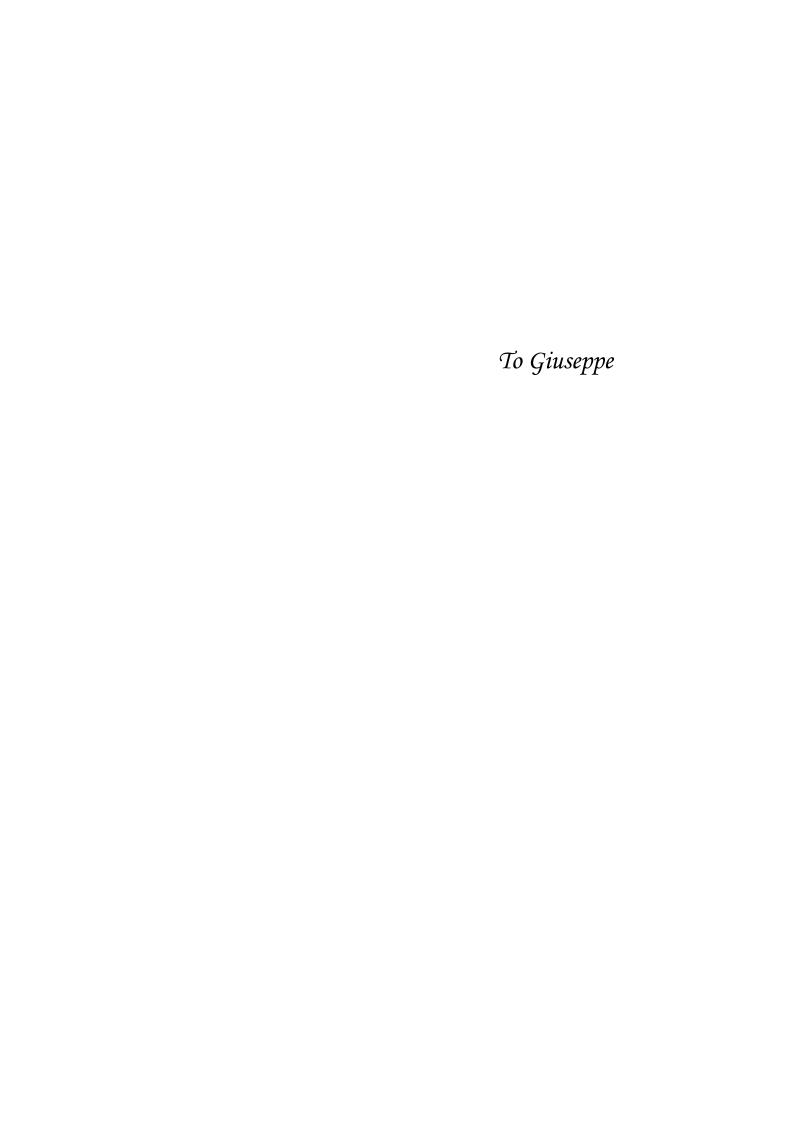
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Notes

The work described in this Thesis was carried out at the International School for Advanced Studies, Trieste, between November 2006 and October 2010.

The work described in this Thesis is included in:

Calcium-activated chloride currents in olfactory sensory neurons from mice lacking bestrophin-2.

Pifferi S, Dibattista M, Sagheddu C, Boccaccio A, Al Qteishat A, Ghirardi F,

Tirindelli R, Menini A.

The Journal of Physiology. 2009 587, 4265-79

I performed caged compounds photolysis patch clamp experiments.

Calcium concentration jumps reveal dynamic ion selectivity of calcium-activated chloride currents in mouse olfactory sensory neurons and TMEM16b-transfected HEK 293T cells.

Sagheddu C, Boccaccio A, Dibattista M, Montani G, Tirindelli R, Menini A.

The Journal of Physiology. 2010 588, 4189–4204

I performed most caged compounds photolysis patch clamp experiments.

ABSTRACT

Olfactory sensory neurons (OSNs) use a Ca²⁺-activated Cl⁻ channels amplification mechanism in olfactory transduction. Odor binding to odorant receptors in the cilia of OSNs leads to an increase of intraciliary Ca²⁺ concentration by Ca²⁺ entry through cyclic nucleotide-gated channels. Ca²⁺ activates a Cl⁻ channel that leads to an efflux of Cl⁻ from the cilia, contributing to the depolarization in OSNs.

The molecular identity of the olfactory Ca²⁺-activated Cl⁻ channel is not definitely established. Bestrophin2 and TMEM16b/anoctamin2 are located at the surface of the olfactory epithelium, in the cilia of OSNs where olfactory transduction takes place.

Moreover when expressed in heterologous systems each of these proteins produces Ca²⁺-activated Cl⁻ currents. Both proteins have been indicated as a candidate for being a molecular component of the olfactory Ca²⁺-activated Cl⁻ channel.

In the first part of this Thesis we analyzed knockout (KO) mice for bestrophin2. We compared the electrophysiological properties of Ca²⁺-activated Cl⁻ currents in OSNs from WT and KO mice for bestrophin2. Our data show that Ca²⁺-activated Cl⁻ currents are still present in the cilia of OSNs from KO mice for bestrophin2 and that their properties are not significantly different from those of WT mice. These results indicate that bestrophin2 does not appear to be the main molecular component of the olfactory Ca²⁺-activated Cl⁻ channel. Therefore further studies are required to determine the physiological function of the bestrophin2 in the cilia of OSNs.

In the second part of this Thesis we measured functional properties of the native Ca²⁺-activated Cl⁻ current in mouse OSNs and compared them with those of TMEM16b/anoctamin2-induced current in transfected HEK cells. We found a similar extracellular blocking potency for some Cl⁻ channels blockers, a similar anion permeability sequence and a reversal potential time-dependency. Therefore, we conclude that the measured electrophysiological properties are largely similar and further indicate that TMEM16b/anoctamin2 is likely to be a major subunit of the native olfactory Ca²⁺-activated Cl⁻ current.

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Abbreviations

AC adenylyl cyclase

ANO anoctamin

BCMCM [6,7-bis (carboxymethoxy) coumarin-4-yl]methyl

BVMD Best vitelliform macular dystrophy

CNG cyclic nucleotide-gated

DPC diphenylamine-2-carboxylate

DIDS 4,4'diisothiocyanostilbene-2,2'-disulphonic acid

DRG dorsal root ganglion

EOG electroolfactogram

FFA flufenamic acid

GPCR G-protein coupled receptor

HEK human embryonic kidney

MeS methanesulfonate

MFA mefenamic acid

NFA niflumic acid

NPPB 5-nitro-2-(3-phenylpropylamino)benzoic acid

OR odorant receptor

OSN olfactory sensory neuron

PDE phosphodiesterase

RPE retinal pigment epithelium

SCN isothiocyanate

SITS 4-acetamido-4'-isothiocyanostilbene- 2.2'-disulphonic acid

TMD transmembrane domain

TMEM transmembrane

1 INTRODUCTION

1.1 Organization of the olfactory system

Chemical senses are responsible for detecting molecules of immense chemical variety from the environment thereby processing specific information concerning food or toxic substances as well as suitable mating partners or predators (reviewed by Firestein, 2001; Tirindelli *et al.*, 2009).

The vertebrate olfactory system is composed of a number of subsystems anatomically segregated within the nasal cavity (reviewed by Breer *et al.*, 2006; Schoenfeld & Cleland, 2005; Ma, 2007; Munger *et al.*, 2009) some well known and others only recently characterized: the main olfactory epithelium, the vomeronasal organ, the Grüneberg ganglion, the septal organ and guanylate cyclase D-containing cells in the main olfactory epithelium (Figure 1.1A). These subsystems are clearly distinguished by the chemosensory receptors they express and the signaling mechanisms they employ to detect and transduce chemosensory stimuli (reviewed by Mombaerts, 1999; Frings, 2001; Breer *et al.*, 2006; Tirindelli *et al.*, 2009); moreover they make distinct neural connections to regions of the olfactory forebrain for processing (reviewed by Munger *et al.*, 2009).

1.1.1 The olfactory epithelium and the olfactory sensory neurons

In vertebrates the main olfactory epithelium lines cartilaginous lamellae, called turbinates, in the posterior nasal cavity. It is a columnar pseudo-stratified neuroepithelium which contains some millions of ciliated olfactory sensory neurons (OSNs), microvillar cells, supporting/sustentacular cells, and basal stem cells (Figure 1.1B) (reviewed by

Breer et al., 2006; Munger et al., 2009; Pifferi et al., 2009c). A layer of mucus is secreted by Bowman's glands at the surface of the olfactory epithelium, partially protecting the tissue that is in direct contact with the external environment. The olfactory epithelium has evolved a constitutive mechanism of neurogenesis (Schwob, 2002) to replenish neuronal population lost during natural turnover or after lesions. At the basal germinal layer, globose and horizontal basal cells are thought to contribute to ongoing neurogenesis but the process is poorly understood (Caggiano et al., 1994; Calof et al., 2002; Leung et al., 2007).

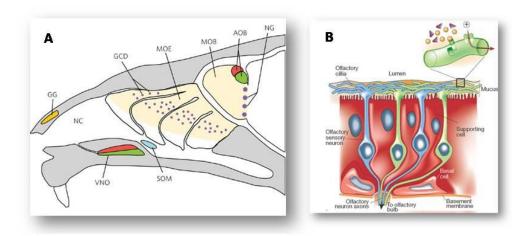


Figure 1.1 A) Cartoon of olfactory sensory system within the nasal cavity of mouse (From Brennan & Zufall, 2006). MOE, main olfactory epithelium; VNO, vomeronasal organ; GG, Grüneberg ganglion; SOM, septal organ of Masera; GCD, guanylate cyclase D-containing cells in the MOE; NC, nasal cavity; NG, necklace glomeruli; MOB, main olfactory bulb; AOB, accessory olfactory bulb; B) Cartoon of the olfactory sensory epithelium primary cells: olfactory sensory neurons, supporting or sustentacular cells and basal stem-cells. Enlargement shows the odor binding at OSNs cilia level (From Firestein, 2001).

Olfactory sensory neurons in mammals are bipolar neurons with the cell body diameter of 5-10 µm (Schild & Restrepo, 1998). The dendrite reaches up to the surface of the tissue with a knob-like ending (diameter 2-3 µm) and several nonmotile cilia (Menco, 1980; Lidow & Menco, 1984; Getchell, 1986; Menco, 1997; Schild & Restrepo, 1998) (Figure 1.2). Cilia are the site of the sensory transduction apparatus (reviewed by Schild & Restrepo, 1998; Nakamura, 2000; Frings, 2001; Kleene, 2008; Pifferi *et al.*, 2009*c*),

their number (20-30) and length (15–50 µm with ~200 nm diameter) increase the cell surface for odor binding (Adamek *et al.*, 1984; Getchell, 1986; Lancet, 1986; Schild & Restrepo, 1998; Kleene, 2008). The axonal processes of OSNs cross the cribriform plate and projects to glomeruli in the olfactory bulb in the most rostral part of central nervous system (Malnic *et al.*, 1999; Mori *et al.*, 1999; Firestein, 2001).

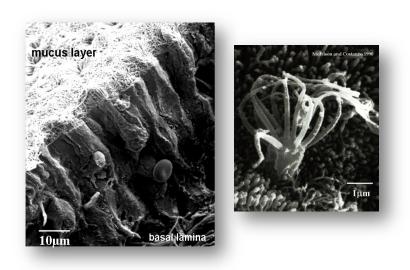


Figure 1.2 Microphotograph of human olfactory epithelium and olfactory sensory neurons knob/cilia obtained with scanning electron microscopy (From Morrison & Costanzo, 1990).

1.2 Odor-induced electrical response in olfactory sensory neurons

In 1955 Ottoson published the first analysis of the electrical activity of the olfactory epithelium by electro-olfactogram (EOG) method (Ottoson, 1955). In EOG recording the odor induced potential change is measured on the surface of the olfactory epithelium (Ottoson, 1955; Scott & Scott-Johnson, 2002; Cygnar *et al.*, 2010).

In isolated OSNs, the response to odor stimuli in solution has been well characterized (Kleene, 2008). Most often, the response has been measured under voltage-clamp upon presentation of a brief pulse of odor (Firestein & Werblin, 1989; reviewed by Kleene, 2008). A given neuron responds to a small and unpredictable subset of odors (Grosmaitre

et al., 2006; Zhao et al., 1998; Lagostena & Menini, 2003). Some evidences suggest that more neurons respond to odors when the epithelium is intact (Sicard & Holley, 1984; Duchamp-Viret et al., 2000) whereas success rates are lower in isolated neurons, typically ranging from 3% to 12% in mouse (Lagostena & Menini, 2003; Reisert et al., 2007).

By directing the odor stimulus to various parts of the cell, it has been shown that the sensitivity to odors is largely restricted to the cilia (Kurahashi, 1989; Firestein *et al.*, 1990; Lowe & Gold, 1993; Takeuchi & Kurahashi, 2003) The odor stimulation generates a transient inward receptor current that would be expected to depolarize the neuron *in situ*. The response typically lasts 1 s or more (Takeuchi *et al.*, 2003; reviewed by Kleene, 2008). In mouse and rat, the latency between arrival of the stimulus and the onset of the current is shorter (~160 ms; Reisert & Matthews, 2001; Grosmaitre *et al.*, 2006) than in amphibians (150 to 600 ms; Firestein *et al.*, 1993; Takeuchi & Kurahashi, 2003; Kurahashi, 1989; Firestein & Werblin, 1987). The amplitude of the peak of the receptor current increases proportionally to concentration or duration of the odor stimulus pulse (Takeuchi & Kurahashi, 2002). The relation between odor dose and peak receptor current is generally well fitted by a Hill equation:

$$I = I_{\text{max}} \frac{C^{n_{\text{H}}}}{C^{n_{\text{H}}} + K_{1/2}^{n_{\text{H}}}}$$

where I_{max} is the maximum macroscopic current, C is the concentration of odor, $K_{\frac{1}{2}}$ is the half-maximally effective concentration, and n_H is the Hill coefficient (Hille, 2001; Kleene, 2008). For some odors $K_{\frac{1}{2}}$ ranges from few to hundred μM both in amphibian (Firestein *et al.*, 1993) and mouse OSNs (Grosmaitre *et al.*, 2006), however some studies report OSN response even at nM and pM odor concentration (Frings & Lindemann, 1990; Grosmaitre *et al.*, 2006).

The parameter $n_{\rm H}$ in Hill equation describes the slope of the dose–response relation. As $n_{\rm H}$ decreases, the slope also decreases, and the dynamic range (range of stimulus strengths over which the neuron response varies) increases. In isolated amphibian OSNs under whole-cell recording conditions, $n_{\rm H}$ ranges from 3-10 (Firestein *et al.*, 1993; Takeuchi & Kurahashi, 2005; Tomaru & Kurahashi, 2005), whereas with suction electrode recordings

or perforated patch analysis, $n_{\rm H}$ is much smaller, 1 to 2 (Ma *et al.*, 1999; Reisert & Matthews, 1999; Grosmaitre *et al.*, 2006).

In physiological solutions, the current-voltage relation of the odor-induced current is nearly linear with a slight outward rectification (Takeuchi & Kurahashi, 2003). The reversal potential is about 0-3 mV (Lowe & Gold, 1993; Takeuchi & Kurahashi, 2003).

1.3 Olfactory transduction

The olfactory transduction of several OSNs involves a canonical cAMP signaling pathway (Figure 1.3) (reviewed by Kleene, 2008; Pifferi *et al.*, 2009*c*). Binding of odorants activates the odorant receptor, which stimulates the rapid synthesis of cAMP by ACIII through a mechanism mediated by the olfaction-specific G protein, $G_{\alpha olf}$.

Cyclic nucleotide-gated (CNG) channels located in the ciliary membrane are directly activated by cAMP, causing a depolarizing influx of Na⁺ and Ca²⁺ ions (Nakamura & Gold, 1987). The increase of the intracellular Ca²⁺ concentration generated by Ca²⁺ entry through CNG channels directly gates Ca²⁺-activated Cl⁻ channels (Kleene & Gesteland, 1991; Lowe & Gold, 1993; Kurahashi & Yau, 1993; Kleene, 1993*b*).

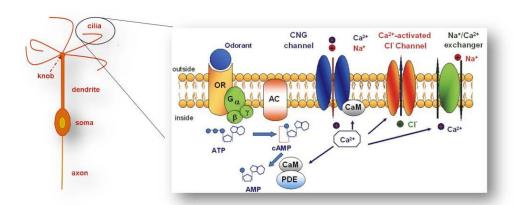


Figure 1.3 Cartoon of olfactory sensory neuron (OSN) and olfactory transduction. The binding of odorant molecules to ORs induces the G protein-mediated activation of ACIII. cAMP directly gates CNG channels generating a depolarizing influx of Na⁺ and Ca²⁺. Ca²⁺ opens a Cl⁻ channel that produces a depolarizing efflux of Cl⁻ (From Pifferi et al. 2006a).

In OSNs the Na⁺/Ca²⁺/2Cl⁻ cotransporter NKCC1 (Kaneko *et al.*, 2004; Reisert *et al.*, 2005; Nickell *et al.*, 2006, 2007; Hengl *et al.*, 2010) and probably the Cl⁻/HCO₃⁻ exchanger SLC4A1 (Hengl *et al.*, 2010) maintain an unusually high internal concentration of Cl⁻ that is in the same range of the Cl⁻ concentration present in the mucus at the external side of the ciliary membrane (Reuter *et al.*, 1998; Kaneko *et al.*, 2001; Kaneko *et al.*, 2004; Nakamura *et al.*, 1997). Therefore in physiological conditions, the opening of Ca²⁺-activated Cl⁻ channels causes an efflux of Cl⁻ ions from the cilia, corresponding to an inward current that further contributes to the depolarization of OSNs (Kurahashi & Yau, 1993; Lowe & Gold, 1993; Kleene & Gesteland, 1991; Kleene, 1993).

Odor response recovery includes reestablishment of the resting ionic gradients in cilia. A Na⁺/Ca²⁺ exchanger (Noé *et al.*, 1997; Reisert & Matthews, 1998) and a plasma membrane Ca²⁺-ATPase (Antolin *et al.*, 2010) contribute to Ca²⁺ efflux as the odor response terminates (Frings, 2001).

In the next paragraphs the olfactory transduction steps will be discussed in detail.

1.3.1 Odorant receptors

Odorant receptor (OR) proteins belong to the G protein-coupled receptors (GPCRs) family (Buck & Axel, 1991). The number of genes encoding ORs varies considerably among species (reviewed by Mombaerts, 2001; Nei *et al.*, 2008), but they probably represent the largest genes family in the mammalian genome (reviewed by Kaupp, 2010). Most mammals have between 600 and 1,300 OR genes, 12 to 50% of them non-functional pseudogenes (reviewed by Nei *et al.*, 2008). Some ORs with unknown function are also expressed in other cell types and body regions, notably in the kidneys and sperm (Spehr *et al.*, 2003; Pluznick *et al.*, 2009).

OR proteins share common motifs but there is a region of hypervariability in the third, fourth and fifth transmembrane regions (reviewed by Mombaerts, 1999; 2004) thought to be a probable binding site for ligands (reviewed by Nakamura, 2000). Reconstruction of odorant-binding sites would help understanding the molecular mechanisms underlying specificity.

A given OR is expressed only in a small number of olfactory sensory cells (Ressler *et al.*, 1993) and the idea that expression of ORs in OSNs follows the so-called one receptorone neuron rule is widely accepted but has not been proved. It is better established that each neuron expresses only one of the two alleles of a given OR gene (reviewed by Mombaerts, 2004), with a transcription process that is not yet fully understood (Rodriguez, 2007). The dose–response relationship for a given odor varies considerably among neurons (Firestein *et al.*, 1993; Grosmaitre *et al.*, 2006; reviewed by Kleene, 2008) and different odorants are recognized by unique but overlapping ensembles of ORs (reviewed by Kaupp, 2010).

1.3.2 The G protein

Odorant receptors, like other GPCRs, interact with a G_s protein which is able to stimulate adenylyl cyclase enzymatic activity. The receptor interacts with α subunit of G-protein (G_{α}) , GDP on this subunit is exchanged to GTP, and the $\beta\gamma$ -subunit $(G_{\beta\gamma})$ is released from G_{α} allowing G_{α} to activate adenylyl cyclase (reviewed by Nakamura, 2000; Pifferi *et al.*, 2009*c*). Jones & Reed (1989) reported that OSNs express a new variant of the G_{α} , named $G_{\alpha olf}$, localized in olfactory cilia (Menco, 1992). $G_{\alpha olf}$ importance in olfactory transduction has been definitely established by showing that knockout mice for $G_{\alpha olf}$ are anosmic (Belluscio *et al.*, 1998).

1.3.3 The adenylyl cyclase

Odor-induced adenylyl cyclase activity has been shown to depend on GTP presence in olfactory cilia (Pace *et al.*, 1985; Sklar *et al.*, 1986; Shirley *et al.*, 1986).

Adenylyl cyclase found in OSNs by PCR screening (Bakalyar & Reed, 1990) was distinct from before-known AC type I or II, therefore it was named type III (ACIII) (reviewed by Frings, 2001; Nakamura, 2000).

Knockout mice for ACIII are completely anosmic (Wong *et al.*, 2000) supporting the idea that cAMP signaling constitutes the main odor transduction mechanism in OSNs.

1.3.4 The phosphodiesterase

Two phosphodiesterases (PDEs) with distinct cellular localization have been found in OSNs: PDE1C is a Ca²⁺/calmodulin-stimulated PDE (Yan *et al.*, 1995) mainly enriched in the cilia and dendritic knob (Borisy *et al.*, 1992; Yan *et al.*, 1995; Yan *et al.*, 1996; Menco, 2005) while PDE4A is Ca²⁺ insensitive (Conti & Beavo, 2007) and it is present throughout the cell, but is absent from the cilia (Cherry & Davis, 1995; Juilfs *et al.*, 1997; reviewed by Nakamura, 2000). PDE1C was hypothesized to be critical for rapid termination of the OSN response due to its ciliary localization and Ca²⁺ dependency, while PDE4A was not expected to affect OSN responses as it is excluded from the cilia.

Cygnar & Zhao (2009) showed that double knockout $Pde1c^{-/-}$ - $Pde4a^{-/-}$ mice and not single knockout mice for PDE1C or PDE4A displayed response for odorants with reduced amplitude, prolonged termination, and slower onset kinetic compared to the wild type, in EOG recordings. Their data indicate that PDE1C and PDE4A are both necessary and that removal of cAMP from the cilia is substantially impaired when all PDE activity is eliminated, confirming a previous study by Firestein *et al.* (1991).

1.3.5 The cyclic nucleotide-gated channel

In 1987 Nakamura and Gold showed the presence of a cAMP-gated current in excised patches from olfactory cilia of toad. Such a current was then described in many other species including salamander, frog, newt, rat and mouse (Kurahashi & Kaneko, 1991; Firestein *et al.*, 1991; Frings *et al.*, 1992; Kleene, 1994).

CNG channels are activated by the direct binding of cyclic nucleotides to a large C-terminal cyclic nucleotide-binding domain (reviewed by Pifferi *et al.*, 2006*a*). The gating of CNG channels is not very voltage-dependent (Kaupp & Seifert, 2002).

Olfactory CNG channel affinity for cGMP is higher than cAMP: K_{1/2} for cAMP is 3 μM in mouse (Michalakis *et al.*, 2006; Song *et al.*, 2008; reviewed by Pifferi *et al.*, 2006*a*), 4.1 μM in rat (Bönigk *et al.*, 1999) and 2 μM in frog (Kleene, 1999); K_{1/2} for cGMP is 2 μM (reviewed by Nakamura, 2000; Pifferi *et al.*, 2006*a*). The Hill coefficient ranges from

1.3 and 2.3 suggesting that at least 2 molecules of cAMP must bind the channel for gating (reviewed by Kleene, 2008).

With low concentrations of divalent cations on both sides of the membrane, the CNG single channel conductance varies from 8 to 46 pS (Kurahashi & Kaneko, 1991; Larsson *et al.*, 1997; Zufall & Firestein, 1993; Frings *et al.*, 1992; Zufall *et al.*, 1991; reviewed by Kleene, 2008) and the current-voltage relation is almost linear with a slight outward rectification (Bönigk *et al.*, 1999; Kurahashi, 1990; Frings *et al.*, 1992; Kleene, 1993*a*; reviewed by Kleene, 2008).

CNG channels are permeant to all monovalent alkali cations, Na⁺, K⁺, Li⁺, Rb⁺ and Cs⁺ with similar permeability ratios in rat (Frings *et al.*, 1992) and in newt (Kurahashi, 1990). Extracellular divalent cations like Ca²⁺ and Mg²⁺ are permeant but also block this channel at negative potentials (Nakamura & Gold, 1987; Zufall & Firestein 1993; Kleene, 1995) resulting in a single channel conductance from 0.56 to 1.5 pS (Zufall & Firestein, 1993; Kleene, 1997). A complex of Ca²⁺ and calmodulin at the intracellular side lowers the affinity for cAMP (Liu *et al.*, 1994; Chen & Yau, 1994), resulting in a lowering of its open probability (reviewed by Nakamura, 2000).

Leinders-Zufall *et al.* (1997, 1998) demonstrated that during odorants application resting Ca²⁺ concentration in the cilium increases from 40 nM to 300 nM exclusively from Ca²⁺ entry through the CNG channel (reviewed by Nakamura, 2000).

The first cyclic nucleotide-gated channel was cloned in retinal rods (Kaupp *et al.*, 1989), then in OSNs (Dhallan *et al.*, 1990; Ludwig *et al.*, 1990). Nowadays six CNG channel genes have been identified in mammals (Kaupp & Seifert, 2002), four subunits types A and two subunits types B (reviewed by Pifferi *et al.*, 2006a).

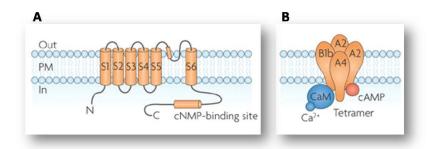


Figure 1.4 Topology (A) and oligomeric state (B) of cyclic nucleotide-gated channel (CNG). Olfactory CNG is composed by one B1b, one A4 and two A2 subunits (From Kaupp, 2010).

Olfactory CNG channels consist of two CNGA2, one CNGA4 and one CNGB1 (b splice variant) subunits (Bradley *et al.*, 1994; Liman & Buck, 1994; Sautter *et al.*, 1998) (figure 1.4). CNGA2 knockout mice lack EOG responses to most odorants (Brunet *et al.*, 1996), though showing residual responses to some other odorants (Zhao & Reed, 2001; Lin *et al.*, 2004).

CNG channels are composed of four subunits forming a tetramer with a central pore (figure 1.4). The topology of each subunit is similar to that of the cationic voltage-activated channels with six transmembrane-spanning domains, a pore-loop domain between the fifth and sixth transmembrane domain, and intracellular N- and C-terminal regions (figure 1.4) (reviewed by Pifferi *et al.*, 2006*a*).

1.3.6 Chloride accumulation in olfactory cilia

In cryosections of rat olfactory epithelium Reuter *et al.* (1998) used energy dispersive X-ray micro analysis and estimated the inner Cl⁻ concentration to be 69 mM and the olfactory mucus Cl⁻ concentration to be 55 mM, with a calculated equilibrium potential for Cl⁻ of +6 mV. In intact olfactory epithelium from mice and rats, Kaneko *et al.* (2004) used two-photon fluorescence lifetime imaging microscopy of the Cl⁻ sensitive dye 6-methoxy-quinolyl acetoethyl ester to measure Cl⁻ concentration in different compartments. They found 50 mM extracellular Cl⁻ and 40-50 mM Cl⁻ concentration in dendritic knobs. Cl⁻ concentration was less in OSN soma suggesting a gradient for Cl⁻ accumulation.

Cl⁻ accumulation process charges the resting cilia to support the excitatory Cl⁻ efflux in olfactory transduction (reviewed by Kleene, 2008) (see paragraph 1.3.7).

The Na⁺/K⁺/2Cl⁻ cotransporter NKCC1 has been shown to be expressed in the cilia of OSNs by proteomic (Stephan *et al.*, 2009; Mayer *et al.*, 2009) and immunological (Hengl *et al.*, 2010) studies. NKCC1 contributes substantially to Cl⁻ uptake (Kaneko *et al.*, 2004) to maintain the intracellular Cl⁻ concentration above electrochemical equilibrium (reviewed by Kleene, 2008). NKCC1 is regulated by phosphorylation of four threonine residues in its N terminus by SPAK (Ste20-related proline-alanine-rich) and OSR1 (oxidative-stress response1) kinases (Dowd & Forbush, 2003; Gagnon *et al.*, 2007;

Delpire & Gagnon, 2008). Both SPAK and OSR1 in turn are regulated by the WNK1 and WNK4 kinases (Anselmo *et al.*, 2006; Delpire, 2009).

Recently Hengl *et al.* (2010) clearly showed ciliary expression of NKCC1 and of the entire regulatory complex for Cl⁻ accumulation in OSNs (see figure 1.5).

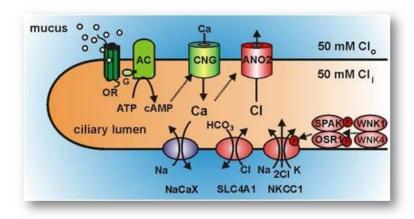


Figure 1.5 Model for signal amplification strategy of olfactory sensory cilia by Hengl et al. (2010). Transduction proteins for this model are in green and amplification proteins in red.

Reisert *et al.* (2005) showed that isolated OSNs do not accumulate Cl⁻ if NKCC1 is eliminated by pharmacological block or genetic removal, but studies in isolated olfactory epithelia (Nickell *et al.*, 2007, 2006) indicated that mice lacking NKCC1 retain approximately 40% to the total Cl⁻ accumulation. Smith *et al.* (2008) also showed normal olfactory sensitivity of NKCC1 knockout mice measured in behavioral test.

The Cl⁻/HCO₃⁻ exchanger SLC4A1 has been shown to localize to the olfactory cilia (Hengl *et al.*, 2010) where it works as an additional mechanism for ciliary Cl⁻ uptake.

1.3.7 The calcium-activated chloride channel

The presence of a Ca²⁺-activated Cl⁻ conductance in OSNs was first discovered in frog olfactory cilia (Kleene & Gesteland, 1991; Kleene, 1993*b*) and it soon became apparent its importance for olfactory transduction. The Ca²⁺-activated Cl⁻ channel

involvement in the response to odorants has been demonstrated in isolated OSNs from amphibians (Kurahashi & Yau, 1993; Firestein & Shepherd, 1995; Zhainazarov & Ache, 1995) and from rats and mice (Lowe & Gold, 1993; Reisert *et al.*, 2005).

Activation of the Ca²⁺-activated Cl⁻ channel is subsequent to the influx of Ca²⁺ through the CNG channel (Kleene, 1993*b*) during odor response.

In voltage-clamp experiments it has been shown that a large fraction of the odor-induced inward current is carried by the secondary Cl⁻ component ranging from 36% in newt and salamander (Kurahashi & Yau, 1993) up to 90% in mice and rats (Lowe & Gold, 1993; Reisert *et al.*, 2005; Nickell *et al.*, 2006; Boccaccio & Menini, 2007) of the total current response.

The current conducted by a single olfactory Cl⁻ channel is very small and single-channel studies have not been possible. By noise analysis of macroscopic currents, the unit conductance was estimated to be 0.8 pS in frog (Larsson *et al.*, 1997), 1.5 pS in rat (Reisert *et al.*, 2003) and 1.6 pS in mouse (Pifferi *et al.*, 2006*b*).

Half maximal activation ($K_{1/2}$) by Ca^{2+} , reported for Cl⁻ conductance, ranges from 2.2 to 4.8 μ M Ca^{2+} (Kleene & Gesteland, 1991; Reisert *et al.*, 2003; Pifferi *et al.*, 2006*b*) showing consistent results for amphibians and rodents. In one other study, $K_{1/2}$ was 26 μ M (Hallani *et al.*, 1998), the reason of such a higher value maybe can be related to the unbuffered Ca^{2+} solution used.

Hill coefficient $n_{\rm H}$ ranges from 2.0 (Kleene & Gesteland, 1991) to 6.6 (Pifferi *et al.* 2006*b*) suggesting that gating of the channel is probably cooperative.

The current-voltage relations in symmetrical Cl⁻ solutions have been shown to rely on Ca^{2+} concentrations: at 2–3 μ M Ca^{2+} , significant outward rectification is apparent, whereas at high Ca^{2+} levels moderate inward rectification is seen (Kleene & Gesteland, 1991; Reisert *et al.*, 2003).

The permeability sequence in inside-out patches are consistent in Reisert *et al.* (2003) ($\Gamma > Br^- > Cl^- > F^-$) and in Pifferi *et al.* (2006*b*) ($\Gamma > NO_3^- > Br^- > Cl^- >> MeS^-$) with "weak field strength" lyotropic series (Wright & Diamond, 1977; Eisenman & Horn, 1983); indicating that the permeability is primarily determined by the hydration energy of the ion. In this model low electrical field strength is associated to the channel binding site, therefore interactions between permeant ions and the channel are weaker than ion-

water interactions. This is why ions with bigger diameter but well dehydrated pass more easily through the channel pore than smaller water-coated ions. Hallani *et al.* (1998) have shown a different ionic permeability sequence ($Cl^- > F^- > I^- > Br^-$), but these data have not been confirmed by other groups.

Niflumic acid (NFA) 300 μM to 1 mM has been shown to reduce by 70-90% the Ca²⁺-activated Cl⁻ current when applied either to the intracellular (Kleene & Gesteland, 1991; Kleene 1993; Reisert *et al.*, 2003; Pifferi *et al.*, 2006*b*) or the extracellular (Lowe & Gold, 1993; Reisert *et al.*, 2005; Pifferi *et al.*, 2006*b*; Boccaccio *et al.*, 2006; Boccaccio & Menini, 2007; Antolin *et al.*, 2010) side of the ciliary membrane.

SITS has been applied at the intracellular side of the olfactory membrane (Kleene & Gesteland, 1991; Pifferi *et al.*, 2006*b*) with a less blocking effect than NFA.

Some other compounds like DIDS, DPC, FFA, have been shown to partially inhibit the olfactory Cl⁻ current (Kleene & Gesteland, 1991), but very specific blockers for Ca²⁺-activated Cl⁻ channels and other Cl⁻ channels are not known.

The presence of a pair of cationic and anionic currents has been thought to be useful to allow depolarizing current responses in a variety of extracellular ionic environments (Kurahashi & Yau, 1993); in fact ciliary membrane of olfactory neurons is exposed to the external environment in the nasal cavity. This is a possible problem for animals such as fish and amphibians whose fresh water habitat could reduce the extracellular Na⁺ concentration and thereby the CNG channels current.

After first studies in amphibians the discovery of the Ca²⁺-activated Cl⁻ current in mammals suggested a more general function for this secondary component in olfactory transduction. Ca²⁺-activated Cl⁻ current plays the key role of high-gain and low-noise amplifier of the primary CNG current (Kleene, 1997) in olfactory transduction.

1.3.8 Calcium clearance in olfactory cilia

In OSNs cilia free Ca²⁺ levels at rest are ~100 nM (Restrepo *et al.*, 1993; Jung *et al.*, 1994; Leinders-Zufall *et al.*, 1997; Saidu *et al.*, 2009), during the odor response free Ca²⁺ concentration increases ranging from 300 nM (Leinders-Zufall *et al.*, 1998) to 100 μ M (Delgado & Bacigalupo, 2004).

The restoration of Ca²⁺ concentrations to pre-stimulus levels (see paragraph 1.3.5) depends on the Na⁺ electrochemical gradient (Reisert & Matthews, 1998; Antolin & Matthews, 2007). Ca²⁺ extrusion mainly occurs via a Na⁺/Ca²⁺ exchanger (NCX) which has been shown to be present in the olfactory dendrite and cilia (Jung *et al.*, 1994; Noé *et al.*, 1997; Castillo *et al.*, 2007; Pyrski *et al.*, 2007). Accumulation of Na⁺ in cilia produced by exchange with Ca²⁺ is likely to be reversed by a Na⁺/K⁺-ATPase which has been identified to be expressed in olfactory cilia (Kern *et al.*, 1991; Lo *et al.*, 1991; Menco *et al.*, 1998; Castillo *et al.*, 2007; Klimmeck *et al.*, 2008; Mayer *et al.*, 2008).

A plasma membrane Ca²⁺-ATPase (PMCA, also called Ca²⁺ pump) has been supposed to further reduce intraciliary Ca²⁺ as the odor response terminates (Lo *et al.*, 1991; Castillo *et al.*, 2007). Ca²⁺-ATPase has been localized to the olfactory cilia by immunological (Weeraratne *et al.*, 2006; Castillo *et al.*, 2007) and proteomic (Klimmeck *et al.*, 2008; Mayer *et al.*, 2008) studies.

The quantitative importance of the Ca²⁺-ATPase appears to be limited compared to the Na⁺/Ca²⁺ exchanger contribution to ciliary Ca²⁺ clearance in OSN both from amphibians (Reisert & Matthews, 1998; Reisert & Matthews, 2001; Antolin & Matthews, 2007; Kleene, 2009; Antolin *et al.*, 2010) and mouse (Saidu *et al.*, 2009).

 Na^+/Ca^{2+} exchanger has a lower affinity for Ca^{2+} with $(K_{\frac{1}{2}} > 1 \ \mu M)$ than Ca^{2+} -ATPase $(K_{\frac{1}{2}} < 0.5 \ \mu M)$ (Carafoli & Brini, 2000). Na^+/Ca^{2+} exchanger could therefore reduce most of the high ciliary Ca^{2+} produced during a strong odor response (see paragraph 1.3.5), Ca^{2+} -ATPase could account for only a small fraction of the Ca^{2+} clearance maybe restoring Ca^{2+} down to the OSN basal level (reviewed by Kleene, 2008).

1.4 Calcium-activated chloride channels in other cell types

Ca²⁺-activated Cl⁻ currents were first described in the 1980s in *Xenopus* oocytes (Barish, 1983; Miledi, 1982) and salamander photoreceptor inner segments (Bader *et al.*, 1982). To date it is well known that Ca²⁺-activated Cl⁻ channels play key roles in several physiological processes including epithelial secretion, membrane excitability in cardiac muscle and neurons, olfactory transduction, regulation of vascular tone (reviewed by Frings *et al.*, 2000; Hartzell *et al.*, 2005).

In oocytes, Ca²⁺-activated Cl⁻ channels play a role in the prevention of polyspermy in amphibians (Webb & Nuccitelli, 1985). In fertilized egg, IP₃ production induces a rapid increase in intracellular Ca²⁺ from internal stores. Ca²⁺ activates Cl⁻ channels resulting in membrane depolarization (reviewed by Hartzell *et al.*, 2005).

In smooth muscle cells, activation of Ca²⁺-activated Cl⁻ channels is part of an amplification mechanism in the regulation of the myogenic tone through membrane depolarization (Leblanc *et al.*, 2005; reviewed by Large & Wang, 1996). Activation of Ca²⁺-activated Cl⁻ channel is mediated by Ca²⁺ entry through the voltage-gated channels or through the Ca²⁺ released from intracellular stores by IP₃ generated in phospholipase C pathway (Large & Wang, 1996; Davis & Hill, 1999). Ca²⁺-activated Cl⁻ channels opening produces a depolarization because chloride equilibrium potential is more positive than the resting potential (Chipperfield & Harper, 2000).

In photoreceptor Ca²⁺-activated Cl⁻ channels are localized in inner segment (Bader *et al.*, 1982; Barnes & Hille, 1989; Maricq & Korenbrot, 1988; Lalonde *et al.*, 2008). The depolarization produced by the dark current opens voltage-gated channels located at the synaptic terminal causing a Ca²⁺ influx that activates a large Cl⁻ conductance. Upon illumination, the dark current turns off, the cell membrane hyperpolarizes, and transmitter release stops (Yau, 1994). The role of Ca²⁺-activated Cl⁻ channel in rods is not known, in cones it has been suggested that this current plays a role in modulating lateral inhibition (Barnes & Hille, 1989; Maricq & Korenbrot, 1988; Thoreson & Burkhardt, 1991).

In airway epithelia activation of Ca²⁺-activated Cl⁻ channels controls the level of mucous hydration which is important for protection against infection. Basally located transporters accumulate Cl⁻ in the cell against the electrochemical gradient. Secretion of fluids is

accomplished by apical Cl⁻ channels that permit Cl⁻ to flow into the extracellular space down its electrochemical gradient (Kunzelmann *et al.*, 2007).

In dorsal root ganglion (DRG) neurons, spinal cord neurons, and neurons of the autonomic nervous system Ca²⁺-activated Cl⁻ channels are thought to regulate neuronal excitability but the mechanism is poorly established (reviewed by Scott *et al.*, 1995). In DRG neurons opening of Ca²⁺-activated Cl⁻ channels by Ca²⁺ entry or Ca²⁺ release from stores would depolarize the cell membrane or produce after-depolarization following action potentials (De Castro *et al.*, 1997; reviewed by Scott *et al.*, 1995; Frings *et al.*, 2000; Hartzell *et al.*, 2005)

Proteins constituting Ca²⁺-activated Cl⁻ channels in most cells and tissues are still elusive. Ca²⁺-activated Cl⁻ currents show heterogeneous biophysical properties, regulatory mechanisms and pharmacology suggesting that different channels are expressed. In many cases, Ca²⁺-activated Cl⁻ channels are activated by Ca²⁺ in a wide range of concentrations (Kuruma & Hartzell, 2000; Hartzell *et al.*, 2005; Angermann *et al.*, 2006) and are also voltage-dependent, with membrane depolarization increasing the activity (Nilius *et al.*, 1997; Lalonde *et al.*, 2008; Hartzell *et al.*, 2005). At not saturating Ca²⁺ concentrations they show outwardly rectifying current-voltage relationship, at maximal Ca²⁺ the current-voltage relationship becomes linear (Kuruma & Hartzell, 2000; reviewed by Frings *et al.*, 2000; Hartzell *et al.*, 2005). In some studies Ca²⁺-activated Cl⁻ currents are directly activated by Ca²⁺, in others, activation requires the intervention of a Ca²⁺/calmodulin-dependent kinase (Arreola *et al.*, 1998; Kaneko *et al.*, 2006).

Despite considerable efforts in the past years these physiologically important ion channels have been difficult to identify at the molecular level. The search for the molecular counterparts for Ca²⁺-activated Cl⁻ currents has been difficult first because a favorite system for expression cloning of ion channels, the *Xenopus* oocyte, is not suitable for these channels since these cells express large endogenous Ca²⁺-activated Cl⁻ currents. Second, drugs to differentiate Ca²⁺-activated Cl⁻ channels from other Cl⁻ channels lack specificity (De La Fuente *et al.*, 2008). Finally, homology cloning has not been fruitful because none of the known cloned Cl⁻ channels have properties that suggest clear structural relationships to Ca²⁺-activated Cl⁻ channels (reviewed by Hartzell *et al.*, 2009).

To date five candidate proteins have been proposed as the molecular counterparts of Ca²⁺-activated Cl⁻ currents: CLCA, ClC-3 and Tweety (reviewed by Hartzell *et al.*, 2005) are briefly described in this section; Bestrophins and TMEM16/Anoctamins (reviewed by Kunzelmann *et al.*, 2009; Hartzell *et al.*, 2009; Flores *et al.*, 2009) will be discussed in detail in the following sections.

CLCA. The Ca²⁺-activated Cl⁻ channel (CLCA) family was cloned from a bovine tracheal cDNA expression library (Ran *et al.*, 1992). Transfect ion of various cell types with cDNAs encoding various CLCAs induces Ca²⁺-dependent currents. However there is great skepticism over the function of CLCAs as chloride channels, because there are too many differences in Ca²⁺ sensitivity, voltage sensitivity and pharmacology with native Ca²⁺-activated Cl⁻ channels (reviewed by Eggermont, 2004; Hartzell *et al.*, 2005), and some of them have very high homology to known cell adhesion proteins and some seem to be soluble, secreted proteins (Gruber & Pauli, 1999).

CIC. The CIC-3 is a member of the CIC family of chloride channels and transporters. The properties of the currents reported for CIC-3, however, differ from those typically described for Ca²⁺-activated Cl⁻ channels (reviewed by Hartzell *et al.*, 2005) and a normal Ca²⁺-activated Cl⁻ channels activity has been shown in parotid acinar cells from a CIC-3 knockout mouse (Arreola *et al.*, 2002).

TWEETY. The human genes hTTYH2 and hTTYH3, with homology to a *Drosophila* gene called tweety, have been shown to encode a Ca²⁺-regulated maxi-Cl⁻ channel (Suzuki & Mizuno, 2004; Suzuki, 2006). Big single channel conductance (260 pS) (Suzuki & Mizuno, 2004) and the absence of this channel in cells with classical Ca²⁺-activated Cl⁻ currents suggest this is not the favorite candidate (reviewed by Hartzell *et al.*, 2005).

1.5 Bestrophins protein family

Proteins of the bestrophin family have been shown to form Cl⁻ channels when expressed in heterologous systems (Sun *et al.*, 2002; Tsunenari *et al.*, 2003) and were proposed to be *bona fide* Ca²⁺-activated Cl⁻ channels (Qu *et al.*, 2004; Qu *et al.*, 2003; Pusch, 2004; Tsunenari *et al.*, 2006; reviewed by Kunzelmann *et al.*, 2007; Hartzell *et al.*, 2008; Marmorstein *et al.*, 2009).

The electrophysiological properties of Cl⁻ channels bestrophins were generally investigated after transient heterologous expression in HEK 293 cells (Sun *et al.*, 2002) (Qu *et al.*, 2003; Qu *et al.*, 2004; Pifferi *et al.*, 2006b) and other cell types.

Different bestrophins produce currents with different current-voltage relations and kinetics of activation. Human bestrophin1 currents have linear current-voltage relationships, essentially time independent, whereas human bestrophin3 currents strongly inwardly rectify and activate slowly with time (Tsunenari *et al.*, 2003).

The ionic permeability of mouse and *Xenopus* bestrophin2 showed a weak selectivity among various anions with the following permeability sequence: $SCN^- > I^- > Br^- > Cl^- > F^-$ (Qu *et al.*, 2003; Qu *et al.*, 2004; Pifferi *et al.*, 2006b), as reported for many Ca^{2+} -activated Cl^- channels (reviewed by Frings *et al.*, 2000; Hartzell *et al.*, 2005).

It is generally agreed that the selectivity of a channel is determined by the channel pore; point mutations in human bestrophin1 and mouse bestrophin2 produce changes in Cl⁻ channel properties (Pusch, 2004; Qu *et al.*, 2004) such as the ionic permeability and/or the channel gating, supporting strong evidence that bestrophin is responsible for forming the channel. However a new study in airway epithelial cells proposed bestrophins as intracellular store calcium modulator (Barro-Soria *et al.*, 2010; reviewed by Edwards & Kahl, 2010).

 Ca^{2+} sensitivity has been investigated for some bestrophins. Human bestrophin1 is activated by increase of intracellular Ca^{2+} concentration with $K_{1/2} \sim 150$ nM (Fischmeister & Hartzell, 2005); mouse bestrophin2 with $K_{1/2} \sim 200$ nM (Qu *et al.*, 2003; Qu *et al.*, 2004). If native bestrophin channels have the same Ca^{2+} sensitivity as these heterologously expressed channels, bestrophin current must be partially activated at all times, because basal free cytosolic Ca^{2+} is typically around 100 nM. Sun *et al.* (2002)

reported that human bestrophin1 current can be rapidly activated by release of Ca²⁺ from caged Ca²⁺, suggesting a direct activation by Ca²⁺. The Ca²⁺ binding site might be located in the C-terminus immediately after the last transmembrane domain because this region contains a high density of acidic amino acids that could coordinate positively charged Ca²⁺. This Asp-rich domain, indicated as a possible Ca²⁺ sensor for bestrophins (Tsunenari *et al.*, 2006), exhibits some similarity with other Ca²⁺-binding domains like the cytoplasmic Ca²⁺ bowl motif of BK_{Ca} potassium channels (Schreiber & Salkoff, 1997; Bao *et al.*, 2004) and the type 3 (T3) motifs in the C-terminal region of thrombospondins (Carlson *et al.*, 2008).

Because of the significant sequence similarity between the Asp-rich domain and the aforementioned Ca²⁺-binding domains, it is plausible to hypothesize that Ca²⁺ activation of bestrophins could involve, at least in part, Ca²⁺-binding to the Asp-rich domain. Molecular dynamics simulations by Kranjc *et al.* (2009) suggest that at least two Ca²⁺-binding sites could be present in the Asp-rich domain of bestrophins possibly involved in Ca²⁺-dependent activation of the channel. In the same study they show through electrophysiological experiments that mutations predicted by their model within the bestrophin Asp-rich domain have an impact on the function, decreasing the Ca²⁺-activated current amplitude.

Single channel properties of endogenous bestrophin have been measured from *Drosophila* S2 cells (Chien *et al.*, 2006). In inside-out patches *Drosophila* bestrophin1 has a single channel conductance of ~2 pS; such a small single channel conductance is a common feature of many Ca²⁺-activated Cl⁻ channels expressed in many cell types (reviewed by Frings *et al.*, 2000; Hartzell *et al.*, 2005).

The human genome contains four bestrophin paralogs (hBest1-4) (Stöhr *et al.*, 2002; Tsunenari *et al.*, 2003), whereas mice have three paralogs (mBest1-3) and one pseudogene (Krämer *et al.*, 2004; reviewed by Hartzell *et al.*, 2008). The first member of the family to be discovered, human bestrophin1 (hBest1 or VDM2), was identified (Marquardt *et al.*, 1998; Petrukhin *et al.*, 1998) as the site of mutation in Best vitelliform macular dystrophy (BVMD - Best's disease), a dominantly inherited, early onset form of macular degeneration (reviewed by Hartzell *et al.*, 2008; Kunzelmann *et al.*, 2009). It was assumed that BVMD is caused by a defect in the basolateral retinal pigmented epithelium

(RPE) cells (Sun *et al.*, 2002; Hartzell & Qu, 2003), where human bestrophin1 was shown to be localized (Marmorstein *et al.*, 2000; Bakall *et al.*, 2003; Mullins *et al.*, 2007). However mice with the bestrophin1 gene disrupted (mBest^{-/-}) have no retinal pathology and Ca²⁺-activated Cl⁻ current in mouse RPE cells is not changed, suggesting bestrophin1 itself could not function as Cl⁻ channel (Marmorstein *et al.*, 2006).

Experimental data for bestrophin transmembrane topology exist mainly for human bestrophin1, but the high conservation of the predicted transmembrane domains, suggests that all vertebrate bestrophins topologies are similar (reviewed by Hartzell *et al.*, 2008). Bestrophins in all of these species have a conserved N-terminal domain that includes the putative transmembrane regions, and a variable C-terminal domain (reviewed by Hartzell *et al.*, 2008). Two topology models have been proposed for hBest1; according to them, the N- and C-terminal domains of bestrophins would be located at the intracellular side of the membrane and would be connected to five (Tsunenari *et al.*, 2003) or four (Milenkovic *et al.*, 2007) hydrophobic domains forming the channel.

Using mutagenesis and cysteine-accessibility analysis of all amino acids from 69 to 105, Qu and co-workers have shown that TMD2 very likely plays a role in ion selectivity of the mouse bestrophin2 pore. Amino acids in TMD2 closer to the C-terminal (A73C, V78C, S79C, and F80C) react with anionic sulfhydryl reagents more slowly than those closer to the N-terminal suggesting that these residues may form the outer mouth of the channel. This is consistent with the C-terminal end of the putative transmembrane domain being closer to the cytoplasm and deeper in the pore. Several other observations indicate that anionic selectivity is determined by this region of the protein. Replacement of F80 with amino acids of opposite charge had opposite effects on rectification of the current: F80R outwardly rectifies, whereas F80E inwardly rectifies (Qu *et al.*, 2004; Qu *et al.*, 2006). Finally the fact that most of the amino acid substitutions that were made in serine 79 (Qu & Hartzell, 2004) disrupted the channel function in qualitatively similar ways, suggests that this residue plays an important role in the pore (reviewed by Hartzell *et al.*, 2008).

Alternative mechanisms for bestrophin functions have been proposed. Barro-Soria *et al*. (2010) showed that endogenous bestrophin1 primarily resides in the endoplasmic reticulum in airway epithelial cells modulating Ca²⁺ release and uptake from intracellular

stores (reviewed by Edwards & Kahl, 2010). Bestrophin1 has also been demonstrated to be permeable to other anions such as HCO_3^- (Qu & Hartzell, 2008) and γ -aminobutyric acid (GABA) (Lee *et al.*, 2010) showing it could be involved in physiological roles other than chloride flux.

1.5.1 Bestrophin2 as olfactory Ca²⁺-activated Cl⁻ channel

Bestrophin2 is expressed in epithelial cells of airways (Barro-Soria et al., 2008), colon (Barro Soria et al., 2009), kidney (Hennig et al., 2008) in the eye (Zhang et al., 2010) and olfactory ciliary epithelium (Pifferi et al., 2006b; Klimmeck et al., 2009). Qu et al. (2004) showed that bestrophin2 heterologously expressed was localized at the cell surface and induced similar Ca²⁺-activated currents in different cell lines (reviewed by Hartzell et al., 2005). Currents induced by bestrophin2 have little or no rectification, with reversal potential as expected for a Cl⁻ selective current and the same anionic permeability sequence $SCN^- > I^- > Br^- > Cl^- > F^-$ (Qu et al., 2004). Mutations in putative pore domains of bestrophin2 (Qu & Hartzell, 2004) alter the conduction and binding of anions. In wild-type channels, substitution of extracellular Cl with SCN produced a significant decrease in conductance and a shift of the E_{rev} . The reduced conductance in both the inward and outward directions by extracellular SCN suggests that not only is SCN less conductive than Cl, it is also able to block outward movement of Cl (inward current), presumably by binding in the channel pore. With the S79C mutant, in contrast, substitution of SCN for Cl did not reduce the conductance and the E_{rev} shifting was smaller.

Bestrophin2 has been proposed as the putative molecular counterpart of Ca²⁺-activated Cl⁻ channels involved in olfactory transduction (Pifferi *et al.*, 2006*b*). Pifferi *et al.* (2006*b*) found by RT-PCR that Bestrophin2 is expressed in OSNs. Bestrophin2 was detected on the cilia by immunocytochemistry, where it colocalizes with CNGA2, the principal subunit of the olfactory CNG channel that is responsible for the primary transduction current. The biophysical and pharmacological properties of the current induced by heterologous expression of bestrophin2 and those of the native Ca²⁺-activated Cl⁻ current from dendritic knob/cilia of mouse olfactory sensory neurons present many

similarities, including the same anion permeability sequence ($\Gamma > NO_3^- > Br^- > C\Gamma^- > MeS^-$), small estimated single-channel conductance, and the same side-specific blockage by some Cl⁻ channel blockers (Pifferi *et al.*, 2006*b*). The most significant difference between the two currents was found to be their sensitivity to intracellular Ca²⁺. In fact, currents were half-maximal at a Ca²⁺ concentration of 0.4 μ M for bestrophin2, whereas native currents required a higher Ca²⁺ concentration, 4.7 μ M (Pifferi *et al.*, 2006*b*).

Two studies challenged the idea of bestrophin2 role in olfactory transduction. Bakall *et al.* (2008) replaced the first two exons of bestrophin2 with Lac-Z and found expression in colon and ciliary epithelium but not olfactory epithelium. Bestrophin2 knockout mice showed no obvious olfactory deficit, but they show diminished intraocular pressure suggesting that bestrophin2 may be involved in aqueous humor generation.

On the contrary Klimmeck *et al.* (2009) clearly showed that bestrophin2 expression was restricted to the cilia of mature OSN and at all subcellular levels in developing sensory neurons. This group suggested a role for bestrophin2 in the process of neurogenesis, during differentiation and growth of axons and cilia of OSNs.

Bestrophin2 has also been suggested to be a cellular volume-regulated Cl⁻ channel (Fischmeister & Hartzell, 2005) because it is strongly inhibited by hyperosmotic and stimulated by hyposmotic solutions; and a HCO_3 ⁻ channel having a relatively high P_{HCO_3}/P_{Cl} (0.69 \pm 0.4 for hBest2 and 0.63 \pm 0.3 for mBest2; (Qu & Hartzell, 2008; Yu *et al.*, 2010).

1.6 Anoctamins/TMEM16 protein family

The TMEM16 family of genes was first described in vertebrates by bioinformatic analyses (Katoh & Katoh, 2003). TMEM16 proteins are well conserved among eukaryotes; humans and mice have 10 genes named as TMEM16A-K, sharing considerable homology (Galindo & Vacquier, 2005). TMEM16A amino acid sequence identity with TMEM16B is larger than 60%, ~40% for TMEM16C, D, and E; whereas TMEM16F, G, H, J, and K, with 20 – 30% identity, probably represent a more distant subgroup of proteins.

S. cerevisiae's sole TMEM16 homologue, called *Ist2p*, plays a role in salt balance and modulation of intracellular ion concentration (Kim *et al.*, 2005; reviewed by Hartzell *et al.*, 2009). *Drosophila melanogaster* has six paralogues. *Axs* (aberrant x-segregation) is ~35% identical to TMEM16H and K, is localized to the endoplasmic reticulum in early embryos, involved in chromosomal nondisjunction and progression of the meiotic cycle (Zitron & Hawley, 1989; Whyte *et al.*, 1993; Krämer *et al.*, 2003; Galindo & Vacquier, 2005).

All TMEM16 proteins have a similar putative topology, consisting of eight transmembrane segments and intracellular NH₂ and COOH termini; a conserved C-terminal domain of unknown function (DUF590) and a N-linked glycosylation site in the last extracellular loop (except in TMEM16K) (reviewed by Flores *et al.*, 2009; Galietta, 2009; Hartzell *et al.*, 2009; Kunzelmann *et al.*, 2009).

A highly conserved hydrophobic region between TM5 and TM6 that protrudes into the membrane is predicted to form a re-entrant p-loop (except in TMEM16H and K) (Katoh & Katoh, 2004*a,b,c*; Katoh & Katoh, 2005; Galindo & Vacquier, 2005)

It has been proposed that some members of the TMEM16 family of membrane proteins, TMEM16A and B, are calcium-activated chloride channels (Caputo *et al.*, 2008; Yang *et al.*, 2008; Schroeder *et al.*, 2008; Pifferi *et al.*, 2009a; Stephan *et al.*, 2009). Because of their eight transmembrane segments topology and their supposed role in anion transport, TMEM16 proteins were also named 'anoctamins' (Yang *et al.*, 2008); therefore they are also referred as ANO1-10.

Some TMEM16 genes have short splice variant transcripts (Bera *et al.*, 2004; Yang *et al.*, 2008), suggesting the possibility that they could have additional non-channel functions or could fulfill tasks in intracellular compartments (Tsutsumi *et al.*, 2004; Mizuta *et al.*, 2007; Schreiber *et al.*, 2010). Current data suggest that members of the TMEM16 family of proteins are involved in both normal vertebrate development and disease; the existence of multiple TMEM16 paralogs in mice and humans might have evolved to allow tissue-specific expression of proteins with similar functions (Galindo & Vacquier, 2005; Rock & Harfe, 2008; Gritli-Linde *et al.*, 2009).

Several members of the TMEM16 family are overexpressed in different types of cancer (Katoh & Katoh, 2003; Galindo & Vacquier, 2005) and mutation of TMEM16E is

associated with the human disorder gnathodiaphyseal dysplasia (GDD) (Tsutsumi *et al.*, 2004; Mizuta *et al.*, 2007). TMEM16G has been shown to promote adhesion between prostate cancer cells (Das *et al.*, 2007, 2008). Because they are accessible cell surface proteins, TMEM16 proteins are a potential drug target for human diseases (Das *et al.*, 2008; reviewed by Hartzell *et al.*, 2009).

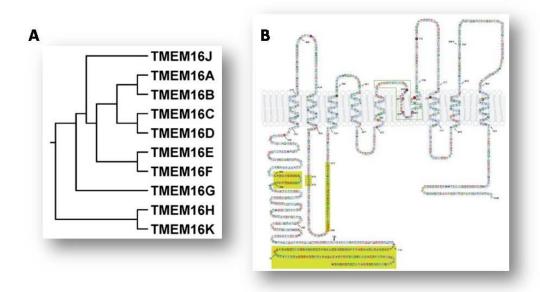


Figure 1.6 A) Phylogenetic tree of human TMEM16 members (From Schroeder et al. 2008). B) TMEM16A protein containing all of its potential alternatively spliced exons (the 'abcd' form, Caputo et al. (2008) Alternatively spliced segments a, b, c and d are shown with a chartreuse background. Single amino acid codes are in circles colored according to the physical properties of the amino acid (green, hydrophilic; pale blue, hydrophobic; red, acidic; magenta, basic; cyan, other ionizable (tyrosine and histidine); yellow, proline; gold, glycine; pink, cysteine) (From Hartzell et al. 2009).

At this point in time, it has been shown experimentally that TMEM16A and B are calcium-activated chloride channels but it is not clear whether it is the same for all TMEM16 members, and TMEM16H has been shown (Schreiber *et al.*, 2010) to inhibit TMEM16A currents. Also, there are no data showing whether different members of the family form heteromers with other members.

1.6.1 TMEM16A/Anoctamin1

TMEM16A is expressed in many of the tissues that are known to express Ca²⁺-activated Cl⁻ currents: airway epithelium and smooth muscle cells, acinar cells of pancreas and salivary glands, proximal kidney tubule epithelium, retina, dorsal root ganglion sensory neurons, and submandibular gland (Huang *et al.*, 2009; Schreiber *et al.*, 2010).

In 2008, three research groups have arrived independently at the identification of TMEM16A as a membrane protein strongly related to the activity of Ca²⁺-activated Cl⁻ channels. TMEM16A expressed in different cell systems, HEK-293, FRT cells and *Axolotl* oocytes always leads to the appearance of currents very similar to classical Ca²⁺-activated Cl⁻ channels (Caputo *et al.*, 2008; Schroeder *et al.*, 2008; Yang *et al.*, 2008).

At nonmaximal Ca²⁺ concentrations, voltage pulses to positive membrane potentials elicit slow-activating currents. This effect is reversible, as the return to negative membrane potentials causes a deactivation of the current. Accordingly, the steady-state current-voltage relationship under this condition is outwardly rectifying.

At maximal concentration of Ca^{2+} , the channels become fully active at all membrane potentials, and consequently, the relaxation after voltage steps disappear (reviewed by Kunzelmann *et al.*, 2009).

Similar to the case for native Ca^{2+} -activated Cl^{-} currents, the Ca^{2+} sensitivity of the channel in excised inside-out patches was voltage dependent. At -60 mV the $K_{\frac{1}{2}}$ for Ca^{2+} was 2.6 μ M, whereas it decreased to 0.4 μ M at +60 mV (Yang *et al.*, 2008).

Schroeder *et al.*, (2008) shows that expression of xTMEM16A in *Axolotl* oocytes produces currents with multiple components with different reversal potentials. These findings suggest that xTMEM16A has multiple open states that differ in their gating kinetics and ionic selectivity.

At present there are no data about the Ca^{2+} binding site: TMEM16A does not have any obvious E-F hand-like Ca^{2+} -binding sites or IQ-domain CaM binding sites. This could mean that either the protein requires another subunit to confer Ca^{2+} sensitivity or that the Ca^{2+} binding site is a novel type that is not easily recognized. If another subunit is required for Ca^{2+} sensitivity, this subunit must be expressed endogenously in the

expression systems used. (reviewed by Flores *et al.*, 2009; Galietta, 2009; Hartzell *et al.*, 2009; Kunzelmann *et al.*, 2009).

TMEM16A has at least four alternatively spliced exons resulting in proteins having between 712 and 1006 amino acids. The alternatively spliced exons include two at the cytoplasmic N-terminus and two in the first cytoplasmic loop (Caputo *et al.*, 2008; Ferrera *et al.*, 2009). All splice variants produced Ca²⁺-activated Cl⁻ currents when expressed in HEK cells, but the variant without any of these segments was not functional (Caputo *et al.*, 2008). This finding suggests that alternative splicing is a mechanism to regulate channel properties and may be the basis for generation of different Ca²⁺-activated Cl⁻ channel types with different voltage dependence and Ca²⁺ sensitivity in a tissue-specific manner (Ferrera *et al.*, 2009).

The region between the fifth and sixth transmembrane segments is predicted to form a reentrant loop important for the formation of the channel pore. Mutagenesis of positively charged amino acids localized in this region (Yang *et al.*, 2008) altered ion selectivity of the channel, thus enhancing its permeability to cations. Altered ion selectivity and voltage dependence were also observed by mutagenesis of an arginine and a glutamine in the third and sixth transmembrane domains, respectively (Caputo *et al.*, 2008).

Another characteristic of TMEM16A is its overexpression in human cancers. Its transcript was also identified as expressed at high levels in gastrointestinal stromal tumors and oral squamous cell carcinomas and therefore also named DOG1 (West *et al.*, 2004) and TAOS2 (Huang *et al.*, 2006). The relationship between cancer and a protein with a role in Cl⁻ transport is not clear. Ca²⁺-activated Cl⁻ channels may be important in proliferation, migration, and resistance of cancer cells to apoptotic stimuli (reviewed by Galietta, 2009). TMEM16A was also found to be one of the candidate genes responsible for autosomal recessive hearing impairment (Kalay *et al.*, 2007). This is of interest because during mouse development TMEM16A is strongly expressed in inner ear cells which will later form the organ of Corti, and parts of the stria vascularis (Gritli-Linde *et al.*, 2009). Finally TMEM16A has an important role in the physiology of airway epithelium and is a possible pharmacological target to circumvent the Cl⁻ transport defect in cystic fibrosis patients.

Rock *et al.* (2008) had reported a TMEM16A knockout mouse. All knockout homozygous mice died within one month of birth showing a severe phenotype characterized by altered formation of tracheal cartilage rings (tracheomalacia). At the moment the reason for tracheal cartilage abnormalities is unknown; it was suggested that the defect in the cartilage rings should be secondary to the improper embryonic stratification of the embryonic tracheal epithelium, which may point out to a functional crosstalk between epithelium and the submucosal tissue during development (reviewed by Flores *et al.* 2009; Galietta, 2009).

Because animals lacking expression of TMEM16A die shortly after birth (Rock *et al.*, 2008), long-term observations are currently not possible.

TMEM16A loss was further analyzed by Ousingsawat *et al.* (2009) in a broad spectrum of epithelial tissues, including airways, colonic epithelium, pancreatic acinar cells, salivary gland cells and hepatocytes, showing TMEM16A contribution to Ca²⁺-activated whole cell currents and to Ca²⁺-dependent Cl⁻ secretion.

Rock *et al.* (2009) found accumulation of mucus in the lumen of tracheas of TMEM16A null mice, suggesting an important function of TMEM16A for mucociliary clearance in mouse airways.

1.6.2 TMEM16B/Anoctamin2

Among the members of the mouse family, TMEM16B is the most similar to TMEM16A, with ~60% amino acid identity (Katoh & Katoh, 2003; Galindo & Vacquier, 2005). TMEM16B transcripts have been found in the retina photoreceptors, in olfactory bulb, olfactory epithelium, pancreas, salivary glands and some brain regions (http://www.brain-map.org).

Large deletions of TMEM16B N-terminus together with von Willebrand factor genes are involved in some cases of the severe von Willebrand disease type 3 (Schneppenheim *et al.*, 2007). Furthermore a recent genome-wide association study (Otowa *et al.*, 2009) in a Japanese population indicated that single nucleotide polymorphisms in TMEM16B gene were significantly associated with panic disorder.

Schroeder *et al.* (2008) and Pifferi *et al.* (2009*a*) reported that TMEM16b generated Ca²⁺-activated Cl⁻ currents in *Axolotl* oocytes and HEK 293 cells.

TMEM16b has been pointed to as a promising candidate for the olfactory Cl⁻ current.

Stephan *et al.* (2009) characterized the mouse olfactory TMEM16b isoform, composed of 24 exons (909 amino acids), with a predicted molecular weight of ~110 kDa. Exon 3, which encodes 33 amino acids in the predicted N-terminal cytoplasmic domain, is lacking in a minority of transcripts in both OSNs and retinal cells, where TMEM16b was first studied (Stöhr *et al.*, 2009). The olfactory TMEM16b variant also lacks the exon 13 (4 amino acids of unknown function) in the first intracellular loop in the retinal variant.

The presence of TMEM16b in OSN has been demonstrated by using different techniques. By in situ hybridization TMEM16b mRNA has been shown to be highly and specifically localized in mature OSNs within the mouse olfactory epithelium (Yu *et al.*, 2005; Hengl *et al.*, 2010; Rasche *et al.*, 2010) and not to the sustentacular or basal stem cell layers.

TMEM16b protein was identified from a proteomic screen of olfactory cilia membrane preparation (Mayer *et al.*, 2009; Stephan *et al.*, 2009; Rasche *et al.*, 2010; Hengl *et al.*, 2010) and by immunocytochemistry it was detected in olfactory epithelium limited to the sensory cilia, where TMEM16b colocalized with marker proteins for olfactory cilia such as acetylated tubulin, CNGA2 (Rasche *et al.*, 2010) and ACIII (Hengl *et al.*, 2010).

Finally in mouse olfactory epithelium infected with adenoviral vector, TMEM16b protein localized primarily in the cilia and dendritic knobs of OSNs (Stephan *et al.*, 2009), demonstrating that TMEM16b is able to use the ciliary targeting machinery.

TMEM16b isoforms from retina (Stöhr *et al.*, 2009; Pifferi *et al.*, 2009*a*) or OSNs (Stephan *et al.*, 2009) expressed in HEK 293 cells produced Ca²⁺-activated Cl⁻ current with functional properties similar to the native olfactory Cl⁻ current when studied both in whole cell configuration (Stöhr *et al.*, 2009; Pifferi *et al.*, 2009*a*); and in inside out patches (Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009), indicating a possible involvement of TMEM16b in the olfactory signal transduction cascade.

TMEM16b Ca²⁺ sensitivity is slightly voltage-dependent, with a $K_{\frac{1}{2}}$ that ranges from 1.8 μ M (Stephan *et al.*, 2009) to 5.1 μ M (Pifferi *et al.*, 2009*a*) at negative potentials and Hill coefficients ~2 (Stephan *et al.*, 2009; Pifferi *et al.*, 2009*a*)

The single channel conductance associated with TMEM16b by noise analysis was estimated to be ~1.2 pS (Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009).

TMEM16b current–voltage relationship is linear (Pifferi *et al.*, 2009*a*) or with inward rectification (Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009) in saturating calcium conditions, whereas it shows pronounced outward rectification after activation at sub-saturating calcium concentrations (Pifferi *et al.*, 2009*a*).

Halide permeability sequence for TMEM16b is $SCN^- > \Gamma > NO_3^- > Br^- > Cl^- > MeS > F^-$ with consistent permeability ratios (P_X/P_{Cl}) in Stephan *et al.* (2009) and Pifferi *et al.* (2009*a*), which are inversely related to the ions hydration energies.

Activation of the TMEM16b channel by different divalent cations: Sr²⁺ efficiently activated a current almost as well as Ca²⁺, Ba²⁺ activated a small current, and no current was observed upon Mg²⁺ application (Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009).

TMEM16b current is blocked ~70-80% by the intracellular (Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009) and extracellular (Pifferi *et al.*, 2009*a*) application of niflumic acid (NFA) in a voltage-independent manner, and the blockage is reversible. Two other fenamates, flufenamic and mefenamic acids (FFA and MFA), as well as NPPB and SITS, only partially blocked the current, whereas DIDS did not have any blocking effect at the intracellular side (Pifferi *et al.*, 2009*a*). NPPB, SITS, and DIDS, produced a 65-80% partial block of both inward and outward currents when applied extracellularly (Pifferi *et al.*, 2009*a*).

The strong similarities between the properties of native Cl⁻ currents (see paragraph 1.3.7) and currents induced by TMEM16b described here, support the hypothesis that TMEM16b is part of the ciliary Cl⁻ channel and may contribute to the excitatory current in olfactory transduction.

2 MATERIALS AND METHODS

2.1 Dissociation of mouse olfactory sensory neurons

Mice were handled in accordance with the Italian Guidelines for the Use of Laboratory Animals (Decreto Legislativo 27/01/1992, no. 116) and European Union guidelines on animal research (No. 86/609/EEC).

Olfactory sensory neurons were dissociated enzymatically from the olfactory epithelium of 1 to 3-months-old C57Black wild type or Bestrophin2^{-/-} mice (see paragraph 2.6). Mice were anesthetized with CO₂ inhalation, decapitated, and then the head was hemisected sagittally along the septum to expose olfactory turbinates. The olfactory epithelium was removed and transferred in 1 ml of zero-divalent mammalian Ringer's solution with 200 µM cysteine and 2 U/ml papain (Sigma, Milano, Italy) for 8-10 minutes at room temperature. The olfactory epithelium was minced with fine forceps. The reaction was stopped by adding 0.5 ml of Ringer's solution with 0.1 mg/ml BSA (bovine serum albumin), 0.3 mg/ml leupeptin, and 0.02 mg/ml of DNAseI (all from Sigma, Milano, Italy). After centrifugation (300 g for 5 min) the cells were resuspended in 1 ml of Ringer's solution and plated on glass coverslips (WPI, Sarasota, FL), coated with poly-L-lysine and concanavalin-A TypeV (Sigma, Milano, Italy). Before use, dissociated olfactory sensory neurons were allowed to settle for 60 min at +4°C. Only olfactory sensory neurons with clearly visible cilia were used for the experiments.

2.2 Cell culture and transfection

HEK 293T cells were grown in DMEM (GIBCO) supplemented with 10% FBS (Sigma, Milano, Italy), 100 U/ml penicillin, and 100 μ g/ml streptomycin (Sigma, Milano, Italy) at 37°C in a humidified CO₂ incubator.

The full-length, dominant olfactory isoform of the mouse TMEM16b/anoctamin2 cloned into the pAdTrack-CMV EGFP-expressing vector (Stratagene, LaJolla, CA, Figure 2.1), provided by professor Haiqing Zhao of the Johns Hopkins University in Baltimore (Stephan *et al.*, 2009), was transfected into HEK 293T cells by using FuGENE 6 reagent (Roche Applied Science, Mannheim, Germany) according to the manufacturer's protocol. Transfected cells were identified by EGFP fluorescence and used for electrophysiological recordings from 24 to 48 hours after transfection.

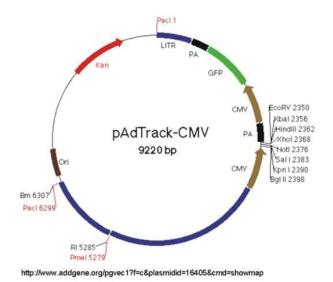


Figure 2.1. pAdTrack-CMV map (www.addgene.org). The full-length, dominant olfactory isoform of the mouse TMEM16b/anoctamin2 was inserted into the multi cloning site (MCS) between NotI and SalI restriction enzymes (Stephan et al., 2009). Transfected HEK 293T cells were identified by EGFP fluorescence and used for electrophysiological recordings.

2.3 The experimental set-up for patch clamp recording

Experiments with olfactory sensory neurons and HEK 293T cells transfected with TMEM16b/anoctamin2 were performed on the same experimental setup, shown in Figure 2.2.

The preparation was observed through an oil immersion 100X objective (N.A. 0.17, Zeiss, Milano, Italy) with an Olympus IX70 inverted microscope (Olympus, Japan) placed on an antivibration table (TMC, USA). A homemade Faraday cage provided adequate electrical shielding.

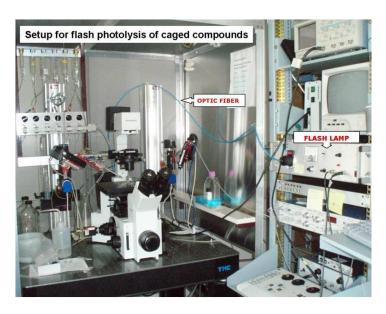


Figure 2.2 Experimental set up used for patch clamp experiments with photolysis of caged compounds.

Patch pipettes were made using borosilicate glass (outer diameter 1.65 mm; inner diameter 1.1 mm, WPI, Sarasota, FL, USA) and pulled with a PP83 puller (Narishige, Tokyo, Japan) using a double-stage pull. The diameter of the tip was about 1 μ m and the pipette resistances were 3–7 M Ω when filled with the standard intracellular solution. Pipettes were mounted in a pipette holder with an Ag/AgCl electrode for electrical recording. The holder movements were controlled by an electronic micromanipulator (Luigs & Neumann, Feinmechanick Elektrotechnik GmbH, Ratingen, Germany).

Currents were recorded in the whole cell voltage-clamp mode with an Axopatch 200B patch-clamp amplifier, controlled by Clampex 8 via a Digidata 1332A (Axon Instruments, Union City, CA, USA). Currents were low-pass filtered at 1 kHz and acquired at 2 kHz for experiments with olfactory sensory neurons, or filtered at 5 kHz and sampled at 10 kHz for experiments with transfected HEK 293T cells.

The perfusion system was entirely gravity driven. Solutions were stored in 50 ml syringes and polyethylene tubes were used for connection with the recording chamber. The recording chamber was continuously bathed with mammalian Ringer solution while an aspiration tube, placed at the opposite site and connected with a trap bottle, controlled the level of solution in the recording chamber. The flow of solution was manually controlled by valves.

In experiments in which the Cl⁻ concentration was changed, the bath was grounded through a 1M KCl agar bridge connected with an Ag/AgCl reference electrode.

All experiments were carried out at room temperature (20 - 24°C).

2.4 Flash photolysis of caged compounds

For flash photolysis of caged compounds a xenon flash-lamp system JML-C2 (Rapp OptoElectronic, Hamburg, Germany) was used coupled with the epifluorescence port of the inverted microscope with a quartz light guide (Boccaccio *et al.*, 2006).

The spot of light had a diameter of about 15 µm to cover only the ciliary region of olfactory sensory neurons (but given the very small size of mouse olfactory sensory neurons it was sometimes technically difficult to restrict the illumination area to the cilia only); or about 50% of the surface of HEK 293T cells (Figure 2.3).

The diameter of the spot was measured by inserting the quartz light guide in an illuminator (Highlight 3100 Olympus) and focusing the spot of the light on the plan of a graduated coverslip. The spot was focused through the oil immersion 100X Zeiss objective used for all the experiments with caged compounds in this thesis.

The released energy at the objective was a few mJ, and this was reduced to an unknown grade of intensity inside the cell.

The flash duration was 1 ms and was kept constant during each experiment. At the beginning of each experiment, the stability of the response was checked by applying repetitive flashes at intervals of about 2 min.

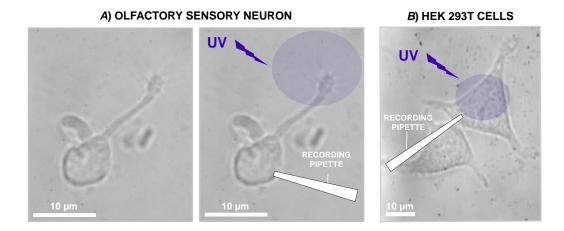


Figure 2.3 Patch-clamp experiments in the voltage-clamp whole-cell configuration were performed on isolated olfactory sensory neuron with clearly visible cilia (A) or HEK293T cells (B). Caged compounds diffused from the patch pipette into the cell and the physiologically active compound was released with ultraviolet light flashes. The illuminated area is indicated by purple circles.

2.5 Ionic solutions

Bath solutions with different ionic composition were used for experiments with olfactory sensory neurons and HEK 293T cells, as listed in the following table:

EXTRACELLULAR SOLUTIONS

	NaCl	KCI	CaCl ₂	MgCl ₂	EGTA	HEPES	Glucose	Na-pyruv.
	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)
Normal Ringer	140	5	1	1		10	10	1
Low Ca ²⁺ Ringer	140	5		1	10	10	10	1

For permeability experiments the NaCl in Normal Ringer was completely replaced with NaSCN,

NaBr, NaI, NaNO₃, NaMeS or CholineCl

All solutions were adjusted to pH 7.4

Niflumic acid (NFA) and 5-Nitro-2-(3-phenylpropylamino)benzoic acid (NPPB, Tocris) were prepared in dimethyl sulfoxide (DMSO) as stock solutions respectively at 200 mM or 83 mM and diluted to the final concentration in the normal Ringer solution (DMSO alone did not alter the currents); 4,4'-diisothiocyanatostilbene-2,2'-disulfonic acid (DIDS) was directly dissolved in the normal Ringer solution. Different bathing solutions were delivered by using a gravity-fed perfusion system. A slow flow rate was selected in such a way that the position of the cilia of the neurons was not perturbed. A complete solution change was obtained in about 10 s. To measure blocker effects, current recordings were obtained before blocker application (control), 1-2 min after delivery of the solution with the blocker, and 2-5 min after perfusion with Ringer solution without the blocker (washout).

In ionic selectivity experiments NaCl was substituted on an equimolar basis with NaX, where X is the substituted anion, or NaCl was replaced with equimolar choline chloride. Relative permeability of the channels was determined by measuring the shift in reversal potential. The bath was grounded through a 1 M KCl agar bridge connected to a Ag/AgCl reference electrode. Liquid junction potentials were calculated using Clampex's Junction Potential Calculator, based on the JPCalc program developed by Barry (1994; see also http://web.med.unsw.edu.au/phbsoft/LJP_Calculator.htm). Applied membrane potentials were corrected off-line. The liquid junction potential between the pipette and the Ringer solution was calculated. Then, if the bathing solution was changed after reaching the

whole-cell configuration, we calculated the additional liquid junction potential generated between the bathing solution and the 1 M KCl agar bridge. We corrected membrane potentials for the following calculated liquid junction potentials (in mV) in the indicated bathing solutions: -4.6 in Ringer, -4.0 in isothiocyanate Ringer, -4.7 in bromide Ringer, -4.6 in iodide Ringer, -4.3 in nitrate Ringer, -3.0 in methanesulfonate Ringer, -5.3 in Ringer with NaCl replaced with choline chloride.

Intracellular recording solutions for the photorelease of 8-Br-cAMP and calcium are summarized in the following table:

INTRACELLULAR SOLUTIONS

	CsCl	KCI	CaCl₂	MgCl ₂	EGTA	HEPES	MgATP	GTP	K-glucon.	BCMCM 8-Br-cAMP	DMNP-EDTA
	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)	(mM)
Caged 8-Br-cAMP		145		4	0.5	10	1	0.1		0.05	
Caged calcium	140		1.5			10					3
Low Cl caged calciun	n	12				10			133		3

All solutions were adjusted to pH 7.4

The caged 8-Br-cAMP (BCMCM-8-Br-cAMP) was provided by Volker Hagen of the Leibniz-Institute for Molecular Pharmacology in Berlin; Boccaccio *et al.*, 2006). The caged 8-Br-cAMP was dissolved in DMSO at 10 or 50 mM and stored at -20° C for up to 3 months. The final concentration of 50 μ M was obtained by diluting an aliquot of the stock solution into the pipette solution.

DMNP-EDTA for the photorelease of caged Ca²⁺ was purchased from Molecular Probes–Invitrogen (West Eugene, OR), and CaCl₂ was adjusted with a 0.1 M standard solution from Fluka (Deisenhofen, Germany).

The caged compounds solutions were stored for a few days at -20° C, kept refrigerated in the dark during the experimental session, and were allowed to diffuse freely from the patch pipette into the cytoplasm of the cell for about 2 min after establishment of the whole cell configuration.

Chemicals, except for caged compounds or otherwise stated, were purchased from Sigma.

2.6 mBest2-null mouse line

The mBest2 null mouse line was purchased from Deltagen (USA). The targeting vector was constructed using 0.8-kb (5') and 2.9-kb (3') mouse Best2 genomic DNA fragments as homology arms. The two arms flanked a promoterless lacZ and a neomycin resistant gene cassette (lacZ-neo). Homologous recombination in mouse embryonic stem cells resulted in the insertion of the lacZ-neo cassette, replacing a region spanning exon 1 through a part of exon 3 of the mouse Best2 locus. Germ-line–transmitting chimeric mice generated from the targeted embryonic stem cells were bred with C57Black mice to produce mBest2^{+/-} mice. Intercrossing of heterozygous mice generated Best2^{-/-} mice.

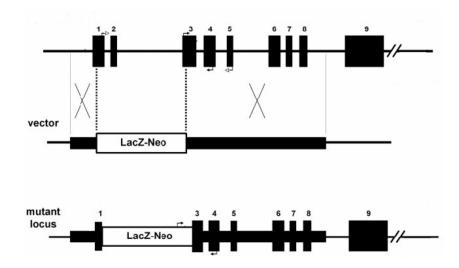


Figure 2. Targeted disruption of the mouse Best2 gene.

Schematic representation of wild-type locus, targeting vector, and mutant locus. Thick lines: fragments used for constructing the targeting vector 5' and 3' arms. Thin lines: genomic DNA or vector backbone sequence. Numbered solid boxes: Best2 exons. Labeled boxes: the LacZ and Neo resistance expression cassette. From Deltagen.

2.7 Data analysis

Current amplitudes at each holding potential were calculated by subtracting the value of the baseline. Data are reported as mean \pm SEM and N indicates the total number of cells. Statistical significance was determined using Student's t-test, or ANOVA, as appropriate. When a statistically significant difference was determined with ANOVA, a Tukey post hoc test was done to evaluate which data groups showed significant differences. P values <0.05 were considered significant.

Data analysis and figures were made with Igor software (Wavemetrics, Lake Oswego, OR, USA).

3 RESULTS

3.1 Calcium-activated chloride currents in olfactory sensory neurons from mice lacking bestrophin-2.

Pifferi S, Dibattista M, Sagheddu C, Boccaccio A, Al Qteishat A, Ghirardi F, Tirindelli R, Menini A (2009). J Physiol. 587, 4265-79

J Physiol 587.17 (2009) pp 4265–4279 4265

Calcium-activated chloride currents in olfactory sensory neurons from mice lacking bestrophin-2

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Olfactory sensory neurons use a chloride-based signal amplification mechanism to detect odorants. The binding of odorants to receptors in the cilia of olfactory sensory neurons activates a transduction cascade that involves the opening of cyclic nucleotide-gated channels and the entry of Ca²⁺ into the cilia. Ca²⁺ activates a Cl⁻ current that produces an efflux of Cl⁻ ions and amplifies the depolarization. The molecular identity of Ca²⁺-activated Cl⁻ channels is still elusive, although some bestrophins have been shown to function as Ca²⁺-activated Cl⁻ channels when expressed in heterologous systems. In the olfactory epithelium, bestrophin-2 (Best2) has been indicated as a candidate for being a molecular component of the olfactory Ca²⁺-activated Cl⁻ channel. In this study, we have analysed mice lacking Best2. We compared the electrophysiological responses of the olfactory epithelium to odorant stimulation, as well as the properties of Ca²⁺-activated Cl⁻ currents in wild-type (WT) and knockout (KO) mice for Best2. Our results confirm that Best2 is expressed in the cilia of olfactory sensory neurons, while odorant responses and Ca²⁺-activated Cl⁻ currents were not significantly different between WT and KO mice. Thus, Best 2 does not appear to be the main molecular component of the olfactory channel. Further studies are required to determine the function of Best2 in the cilia of olfactory sensory neurons.

(Received 1 June 2009; accepted after revision 14 July 2009; first published online 21 July 2009)

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Abbreviations Best2, bestrophin-2; CNG, cyclic nucleotide-gated; KO, knockout; WT, wild-type.

In vertebrates, the process of olfactory transduction occurs in sensory neurons, located in the olfactory epithelium in the nasal cavity. Each olfactory sensory neuron bears several cilia departing from the knob-like swelling of the apical part of the dendrite. The cilia are the site of olfactory transduction: odorant molecules bind to specific receptors expressed in the ciliary plasma membrane activating a G protein-coupled transduction cascade. The activation of adenylyl cyclase by the G protein produces an increase in the ciliary concentration of cAMP, which opens cyclic nucleotide-gated (CNG) channels, which produces a primary inward current carried by Na⁺ and Ca²⁺ ions (reviewed by Schild & Restrepo, 1998; Menini, 1999; Firestein, 2001; Matthews & Reisert, 2003; Menini *et al.* 2004; Pifferi *et al.* 2006*a*; Kleene, 2008). The increase in

While most of the components of the olfactory transduction cascade have been identified at the molecular level, the molecular identity of Ca²⁺-activated Cl⁻ channels is still elusive. In recent years, several proteins have been proposed as possible candidates for Ca²⁺-activated Cl⁻

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Ca²⁺ concentration triggers the gating of Ca²⁺-activated Cl⁻ channels that gives rise to a secondary Cl⁻ current. Since olfactory sensory neurons maintain an elevated intracellular Cl⁻ concentration, which is in the same range of the Cl⁻ concentration present in the mucus at the external side of the cilia (Reuter *et al.* 1998; Kaneko *et al.* 2001, 2004), the opening of Ca²⁺-activated Cl⁻ channels in the ciliary membrane causes an efflux of Cl⁻ ions from the cilia, which amplifies the primary inward current (Kleene & Gesteland, 1991; Kleene, 1993; Kurahashi & Yau, 1993; Lowe & Gold, 1993; Kleene, 1997; Boccaccio & Menini, 2007; reviewed by Frings *et al.* 2000; Kleene, 2008; Frings, 2009).

S. Pifferi and M. Dibattista contributed equally to this study.

channels, including the families of bestrophins, tweety, CLCA calcium activated chloride channels (reviewed by Hartzell *et al.* 2005, 2009) and, very recently, the anoctamin/transmembrane 16 (TMEM16) protein family (Caputo *et al.* 2008; Schroeder *et al.* 2008; Yang *et al.* 2008; Pifferi *et al.* 2009; Stephan *et al.* 2009).

Proteins of the bestrophin family have been shown to form Cl⁻ channels when expressed in heterologous systems (Sun *et al.* 2002; Tsunenari *et al.* 2003) and have been proposed to be *bona fide* Ca²⁺-activated Cl⁻ channels (Qu *et al.* 2003, 2004; Pusch, 2004), although other reports suggested that they function as regulators of ion transport rather than as ion channels (Rosenthal *et al.* 2006; Yu *et al.* 2008; reviewed by Kunzelmann *et al.* 2007; Hartzell *et al.* 2008; Marmorstein *et al.* 2009).

We have previously shown that bestrophin-2 (Best2) is expressed in the cilia of mouse olfactory sensory neurons, where it colocalizes with CNGA2, the principal subunit of the olfactory CNG channel that is responsible for the primary transduction current (Pifferi et al. 2006b). Moreover, we have shown that the functional properties of the current induced by heterologous expression of mouse Best2 and those of the native Ca2+-activated Clcurrent from dendritic knob/cilia of mouse olfactory sensory neurons present many similarities, including the same anion permeability sequence, small estimated single-channel conductances, and the same side-specific blockage by some Cl- channel blockers, although also differences do exist and include a Ca2+ sensitivity discrepancy of one order of magnitude (Pifferi et al. 2006b). However, based on the overall findings, Best2 was indicated to be candidate molecular component of Ca²⁺-activated Cl⁻ channels involved in olfactory transduction (Pifferi et al. 2006b). In the last year knockout mice for Best2 became commercially available opening the possibility of further investigation of the physiological role of Best2. We have therefore analysed the responses of the olfactory epithelium to odorant stimulation and investigated the properties of Ca²⁺-activated Cl⁻ currents in wild-type (WT) and knockout (KO) mice lacking Best2. Our results confirm that Best2 is expressed in the cilia of olfactory sensory neurons, but we found that Ca²⁺-activated Cl⁻ currents were not significantly different between WT and KO mice, indicating that Best2 is not the main molecular component of the olfactory channel. Further studies are required to determine the physiological role of Best2 in the cilia of olfactory sensory neurons.

Methods

Ethical approval

All animals were handled in accordance with the Italian Guidelines for the Use of Laboratory Animals (Decreto

Legislativo 27/01/1992, no. 116) and European Union guidelines on animal research (No. 86/609/EEC). For experiments mice were anaesthetized by CO₂ inhalation and then decapitated.

Animals

Experiments were performed on knockout (KO) mice for *Best2* and wild-type (WT) littermates between 2 and 6 months of age. *Best2* homozygous mutant and WT mice were obtained by breeding heterozygous mutant mice obtained from Deltagen (San Mateo, CA, USA). The generation of these mice has been previously described in detail (Bakall *et al.* 2008).

Cookie test

Mice were left overnight without food with water *ad libitum*. The next day, mice were moved into an opaque cage, while a food pellet (Altromin-R, A. Rieper S.p.A., Vandoies, Bolzano, Italy) was buried in their litter's sawdust, about 2 cm underneath. Then, mice were brought back in their cages and released at the centre of the litter. The time was measured from the moment they were freed to the moment they found the pellet. Results were analysed using the analysis software SPSS 13.0 (SPSS Inc., Chicago, IL, USA) and StatView (SAS Institute Inc., Cary, NC, USA).

RNA isolation and RT-PCR

Total RNA was extracted from the olfactory epithelium of WT and KO mice using standard Clontech procedures (BD Biosciences, Hertfordshire, UK). RNA quality was measured using a NanoDrop1000 Spectrophotometer (ND-1000). Gene expression was examined by RT-PCR from total RNA using primers designed against *Best2*, *CNGA2* and the housekeeping gene *S16*. PCR conditions were as follows: an initial denaturation step of 10 min at 95°C, followed by 35 cycles of 1 min at 94°C, 1 min at 58°C and 1 min at 72°C, and a final extension step of 10 min at 72°C. The products were visualized following agarose gel electrophoresis (1.5%) and DNA was stained with ethidium bromide (10 mg ml⁻¹). Samples without cDNA were used as negative controls. The sequences of the primers used were the following:

Best2 (forward: 5'-AGT CCC AGG AAA CAT AAC AGC TCT C-3' and reverse: 5'-CTC CCA GCA TCT TCC CTT GGC TCA C-3');

CNGA2 (forward: 5'-AGG GAA AGG GCA CCA AAA AGA AA-3' and reverse: 5'-CCA GCA CCA GCC ATA CCA CAA A-3');

S16 (forward: 5'-GGC AGA CCG AGA TGA ATC CTC A-3' and reverse: 5'-CAG GTC CAG GGG TCT TGG TCC-3').

Western blot

Proteins were extracted from olfactory epithelium tissues by RIPA buffer (Millipore, Milan, Italy) and the protein concentration of each sample was determined using the Bio-Rad assay. For Western blotting, 10 µg of protein was separated by gel electrophoresis (SDS-PAG; 12% w/v) and the proteins were electro-blotted onto nitrocellulose filters (Whatman, Germany). Filters were blocked in 1% w/v bovine serum albumin (BSA) in Tris-buffered saline Tween 20 (TBS Tween) and incubated overnight at 4°C with the following primary antibodies: rabbit polyclonal anti-Best2 (1:500; Pifferi et al. 2006b); mouse monoclonal anti-CNGA2 (a gift from F. Müller and U. B. Kaupp, Forschungszentrum Julich, Julich, Germany; Meyer et al. 2000), and anti- β -actin (1:1000; Sigma, Milan, Italy). Membranes were washed in TBS-Tween before staining with antibodies to the appropriate peroxidase-conjugated secondary antibody, diluted 1:1000 in 1% w/v BSA in TBS Tween for 1 h. Blots were developed with the ECL detection system (Amersham, UK).

Immunohistochemistry

The nasal regions were fixed in 4% paraformaldehyde for 4 h at 4°C, decalcified by overnight incubation in 0.5 M EDTA, and then equilibrated in 30% (w/v) sucrose overnight at 4°C for cryoprotection. Coronal sections 16 μ m thick were cut on a cryostat and stored at -20° C. Tissue sections were incubated with 0.5% sodium dodecyl sulfate (v/v) in phospate buffered saline (PBS) for 15 min for antigen retrieval, then incubated in blocking solution (2% normal goat serum, 0.2% Triton X-100 in PBS) for 90 min, and incubated overnight at 4°C in primary antibodies diluted in blocking solution. After rinsing in 0.1% (v/v) Tween 20 in PBS, sections were incubated with fluorophore-conjugated secondary antibodies in 0.1% Tween 20 in PBS for 2 h at room temperature and washed. 4'-6-Diamidino-2-phenylindole (DAPI) $(0.1 \,\mu\mathrm{g}\,\mathrm{ml}^{-1})$ was used to stain nuclei: tissue sections were incubated for 30 min then washed and mounted with Vectashield (Vector Laboratories, Burlingame, CA, USA).

Primary antibodies were rabbit anti-Best2 (Pifferi *et al.* 2006*b*) and mouse monoclonal anti-CNGA2 (Meyer *et al.* 2000) used at 1:50. Secondary antibodies were Alexa 488-conjugated goat anti-rabbit and Alexa 594-conjugated goat anti-mouse diluted to 1:200 (Molecular Probes-Invitrogen, Eugene, OR, USA).

Images were visualized by Leica TCS SP2 confocal microscope, acquired using Leica software at

 1024×1024 pixels resolution and analysed with ImageJ software.

Electro-olfactograms

Electro-olfactogram (EOG) recordings were performed as previously described (Franceschini *et al.* 2009). The mouse head was cut sagitally to expose the medial surface of the olfactory turbinates and EOG recordings were measured at the surface of the olfactory epithelium in response to odorant stimuli in the vapour phase. Each odorant, amylacetate, cineole and acetophenone (Sigma, Milan, Italy), was prepared as 2.5 M stock in DMSO and then diluted with water to the final concentrations used in the experiments. Responses to DMSO alone were less than 0.05 mV. Vapour-phase odorant stimuli were generated by placing 0.9 ml of an odorant solution in a 10 ml glass test-tube capped with a rubber stopper. For stimulation, a 100 ms pulse of the odorant vapour at 8 psi was injected into a continuous stream of humidified air.

Electrophysiological recordings from dissociated olfactory sensory neurons

Olfactory sensory neurons were dissociated enzymatically from the olfactory epithelium of 1- to 2-month-old mice, with a papain-cysteine treatment as previously described (Lagostena & Menini, 2003; Boccaccio *et al.* 2006). Cells were plated on Petri dishes for excised patch recordings, or on glass coverslips coated with poly-L-lysine and concanavalin A (Type V, Sigma, Milan, Italy) for whole-cell recordings with photolysis of caged compounds (Boccaccio *et al.* 2006; Boccaccio & Menini, 2007).

Currents in the whole-cell or in the inside-out voltage-clamp modes were recorded with an Axopatch 1D or an Axopatch 200B amplifier controlled by Clampex 8 or 9 via a Digidata 1322A or 1332A (Axon Instruments, Union City, CA, USA). Patch pipettes were made using borosilicate capillaries (WPI, Sarasota, FL, USA) and pulled with a Narishige PP83 puller (Narishige, Tokyo, Japan). Patch pipettes filled with standard intracellular solutions had resistances of 2–7 $M\Omega$ for whole-cell and 7–10 $M\Omega$ for excised patch recordings. Currents were low-pass filtered at 1 kHz and acquired at 2 kHz for whole-cell experiments, or filtered at 4 kHz and sampled at 10 kHz for excised patch recordings. All experiments were carried out at room temperature (20–22°C).

For flash photolysis of the caged compounds, we used a xenon flash-lamp system, JML-C2 (Rapp OptoElectronic, Hamburg, Germany), coupled with the epifluorescence port of the microscope with a quartz light guide (Boccaccio *et al.* 2006). The spot of light had a diameter of about $15 \,\mu m$ and was focused on the ciliary region. The interval

between experiments was about 2 min to allow the cell to recover from adaptation.

Rapid solution exchange in inside-out patches was obtained with the perfusion Fast-Step SF-77B (Warner Instrument Corp., Hamden, CT, USA). For current–voltage relations of Ca^{2+} -activated currents, inside-out patches were pre-exposed to the test Ca^{2+} concentration for 500 ms at -100 mV to allow the current to partially inactivate, and then a double voltage ramp from -100 to +100 mV and back to -100 mV was applied at 1 mV ms $^{-1}$. The two current–voltage relations were averaged and leak currents measured with the same ramp protocol in Ca^{2+} -free solutions were subtracted. The same type of voltage protocol was used to measure current–voltage relations of cAMP-activated currents.

Ionic solutions

For whole-cell recordings, the extracellular mammalian Ringer solution contained (in mm): 140 NaCl, 5 KCl, 1 CaCl₂, 1 MgCl₂, 10 Hepes, 10 glucose and 1 sodium pyruvate (pH 7.4). The composition of the low Ca²⁺ extracellular solution was similar, except that it contained 10 mm EGTA and no added Ca²⁺. The whole-cell pipette solution for the photorelease of caged 8-Br-cAMP contained (in mm): 145 KCl, 4 MgCl₂, 0.5 EGTA, 10 Hepes, 1 MgATP, 0.1 GTP, 0.05 caged 8-Br-cAMP, (pH 7.4). The caged BCMCM-8-Br-cAMP (Boccaccio et al. 2006) was dissolved in DMSO at 10 or 50 mM and stored at -20° C for up to 3 months. The final concentration of 50 μ M was obtained by diluting an aliquot of the stock solution into the pipette solution, kept refrigerated in the dark during the experimental session, and stored for a few days at -20° C. The standard pipette solution for the photorelease of caged Ca²⁺ contained (in mm): 3 DMNP-EDTA, 1.5 CaCl₂, 140 KCl, and 10 Hepes (pH 7.4). The low Cl⁻ intracellular solutions for the photorelease of caged Ca²⁺ contained (in mm): 3 DMNP-EDTA, 1.5 CaCl₂, 12 KCl, 133 potassium gluconate, and 10 Hepes (pH 7.4). Liquid junction potentials were corrected off-line. DMNP-EDTA was purchased from Molecular Probes-Invitrogen, and CaCl₂ was adjusted with a 0.1 M standard solution from Fluka (Deisenhofen, Germany). The caged compounds were allowed to diffuse from the patch pipette into the cytoplasm of an olfactory sensory neuron for about 2 min after establishment of the whole-cell configuration.

For inside-out recordings, the standard solution in the patch pipette contained (in mm): 140 NaCl, 10 HEDTA and 10 Hepes, pH 7.2. In experiments for Ca²⁺ dose–response relations, NaCl was replaced with LiCl to inhibit the Na⁺/Ca²⁺ exchanger. The bathing solution at the intracellular side of the patch contained (in mm): 140 NaCl or LiCl, 10 HEDTA and 10 Hepes, pH 7.2, and no added Ca²⁺ for the nominally 0 Ca²⁺ solution, or

various added Ca²⁺ concentrations, as calculated with the program WinMAXC (C. Patton), to obtain free Ca²⁺ in the range between 1.5 and 100 μ M (Patton *et al.* 2004). The free Ca²⁺ concentrations were experimentally determined by Fura-4F (Molecular Probes–Invitrogen) measurements by using an LS-50B luminescence spectrophotometer (PerkinElmer, Wellesley, MA, USA). To activate CNG channels a solution containing 100 μ M cAMP directly dissolved into the 0 Ca²⁺ bathing solution was used.

Chemicals, unless otherwise stated, were purchased from Sigma (Milan, Italy).

Data analysis

Data are reported as means \pm standard deviation, with the number of experiments (n) from different mice, cells or membrane patches, as appropriate. The statistical significance of data was evaluated by Student's t-test and P values < 0.05 were considered significant. Data analysis and figures were made with Igor software (Wavemetrics, Lake Oswego, OR, USA).

Results

Expression of Best2 in the olfactory epithelium

To examine the expression of *Best2*, we performed RT-PCR on total RNA from the olfactory epithelium of WT and KO mice. Specific primers for *Best2*, for the main subunit of the CNG channel *Cnga2*, and for the housekeeping gene *S16* showed that, in WT mice, PCR products of the predicted size were amplified (Best2, 205 bp; CNGA2, 200 bp; S16, 102 bp) (Fig. 1A). In KO mice, the 205 bp reaction product, corresponding to *Best2*, was absent, while control genes were normally expressed (Fig. 1A). This result confirms the absence of expression of *Best2* in the olfactory epithelium of KO mice.

Best2 immunoreactivity in the olfactory epithelium

To verify the lack of expression of the Best2 protein in the olfactory epithelium of KO mice, we performed both Western blotting and immunohistochemistry (Fig. 1*B* and *C*), using the antibody against Best2 that we have previously generated and characterized (Pifferi *et al.* 2006*b*).

By Western blotting, we identified a 57 kDa band in a membrane fraction of the olfactory epithelium of WT mice, corresponding to the expected molecular weight for the Best2 protein, in agreement with our previous study (Pifferi *et al.* 2006*b*). The 57 kDa band was undetectable in KO animals, while both the 75 and 42 kDa bands, corresponding respectively to CNGA2 and β -actin, were expressed in both mouse lines (Fig. 1*B*).

By immunohistochemistry, we confirmed our previous results showing that Best2 is expressed in the olfactory epithelium of WT mice (Pifferi *et al.* 2006*b*). We found staining at the surface of the olfactory epithelium, at the level of the ciliary layer, where Best2 colocalized with CNGA2 (Fig. 1*C*, top panels). In the olfactory epithelium of KO mice, Best2 immunoreactivity was absent, while CNGA2 was normally expressed at the level of the ciliary layer (Fig. 1*C*, bottom panels). These results demonstrate the loss of the Best2 protein in the olfactory epithelium of KO mice and confirm the specificity of our antibody against this protein.

Behavioural olfactory response

To determine whether deletion of *Best2* caused a behavioural olfactory deficit, mice were examined for olfactory function in a cookie-finding test. In this test, we compared the food-finding ability of WT and KO mice by measuring the latency to locate buried food. Mice were food-deprived with free access to water overnight and then were put in a cage where a food pellet was buried under the litter. None of the mice searched randomly in the litter; conversely, they dug only in the place where the pellet was hidden. The average time necessary to locate the cookie

was 55 ± 32 s (n = 22) for WT mice, similar to 62 ± 34 s (n = 22) for KO animals. These results show that KO mice do not exhibit any gross olfactory deficit, in agreement with previous results (Bakall *et al.* 2008).

Odorant-induced responses in WT and KO mice

To investigate whether disruption of the Best2 gene modifies the odorant sensitivity of olfactory sensory neurons, we measured odorant-induced changes in voltage across the olfactory epithelium of WT and KO mice. Indeed, the electrical activity of a population of olfactory sensory neurons in response to odorants can be recorded at the surface of the olfactory epithelium as a negative electrical field potential, the electro-olfactogram (EOG) (Ottoson, 1955; Scott & Scott-Johnson, 2002). EOG responses induced by delivering the vapour phase of a 2.5 M amyl acetate solution for 100 ms to the olfactory epithelium were recorded at 13 different locations as indicated in Fig. 2A. Although the amplitudes of EOG responses varied according to the different subregions of the olfactory epithelium, amplitudes at each specific location were not significantly different between WT and KO mice (Fig. 2B). Similar results were obtained with two

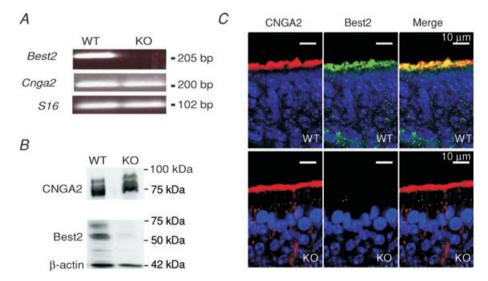


Figure 1. Comparison of Best2 mRNA expression and Best2 immunoreactivity in the mouse olfactory epithelium of WT and KO mice

A, reverse transcription–polymerase chain reaction (RT-PCR) derived cDNA products amplified from RNA of the olfactory epithelium in WT and KO mice using specific primers for *Best2*, *CNGA2* and *S16*, as indicated in the figure. The predicted size of the products for *Best2*, *CNGA2* and *S16* was respectively 205, 200 and 102 base pairs (bp). *B*, Western blot analysis of proteins of the olfactory epithelium in WT and KO mice probed with antibodies against Best2, CNGA2 and β -actin. Bands of the appropriate molecular mass were observed for each protein in WT mice, whereas only bands corresponding to CNGA2 and β -actin were detected in KO mice. The expected molecular mass for Best2, CNGA2 and β -actin was respectively 57, 75 and 42 kDa. *C*, immunostaining of sections of the olfactory epithelium. Confocal micrographs showing Best2 and CNGA2 expression in the ciliary layer of the olfactory epithelium of WT and KO mice. CNGA2 and Best2 co-expression was evident in WT mice, whereas no immunoreactivity to Best2 was detectable in KO mice. Each image on the right was obtained from the merge of the respective left and centre images. Cell nuclei were stained by DAPI.

other commonly used odorants: acetophenone and cineole (data not shown).

We determined the dose–response relation in response to amyl acetate by delivering the vapour phase of odorant solutions at various concentrations to the olfactory epithelium. Figure 2C shows representative recordings of EOG responses to amyl acetate in WT and KO mice. The odorant concentration producing 50% of the maximal EOG amplitude was about 10^{-3} M for both WT and KO mice (Fig. 2D).

We further analysed the kinetics of the EOG recordings. We measured the latency of the response as the interval between the beginning of the odorant application and the time at which the response reached 1% of its maximal value, the rise time as the time interval between 1% and the peak EOG response, and the termination as the time constant of the exponential fit of the recovery phase of the EOG response from the peak value to 10% of the peak. We did not find any significant difference for any of these parameters between WT and KO mice (Fig. 3).

Thus, no differences were observed between KO and WT mice in EOG recordings, indicating that the absence of Best2 does not significantly affect responses to odorants. However, we cannot exclude the possibility that EOG

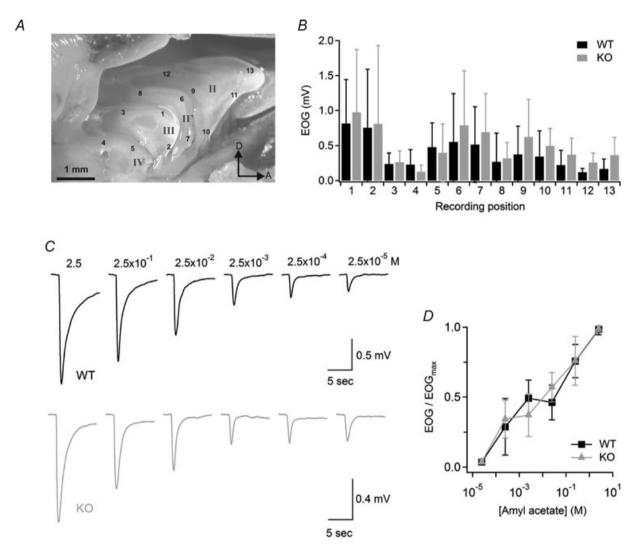


Figure 2. Odorant sensitivity in WT and KO mice

A, photomicrograph of the olfactory turbinate system. Roman numbers designate individual turbinates. Arabic numbers indicate the locations where EOG responses were recorded. D, dorsal; A, anterior. B, average EOG amplitudes in response to a 100 ms pulse of odorant vapour from a bottle containing 2.5 m amyl acetate liquid solution measured at the locations indicated in A (n = 7-14). C, representative EOG recordings from WT (black traces) or KO (grey traces) mice in response to 100 ms pulses of amyl acetate vapours. Numbers above traces are the concentrations of amyl acetate solutions in the bottle. EOG recordings were from location 1. D, EOG amplitudes were normalized to the value measured in response to the vapour of a 2.5 m amyl acetate solution, averaged, and plotted *versus* amyl acetate concentrations in solution for WT (n = 14; black symbols) or KO (n = 13, grey symbols) mice. Data points are linked with straight lines.

recordings are similar in WT and KO animals because some compensatory mechanism may modify the intraciliary ion concentrations in KO compared to WT mice. Indeed, although EOG measurements have the advantage of allowing long recordings while leaving the neurons in a relatively unperturbed situation, this technique does not allow the control of the intracellular ionic composition of neurons and of the membrane potential.

Currents in isolated olfactory sensory neurons

To achieve a control of both the intracellular and extracellular ionic compositions, as well as of voltage, we used isolated olfactory sensory neurons and the patch-clamp technique in the whole-cell voltage-clamp configuration. To investigate whether a Ca²⁺-activated Cl⁻ current was present in individual olfactory sensory neurons from WT and KO mice, we measured the transduction current directly activating CNG channels in the cilia (Fig. 4). Indeed, the use of odorants to activate the transduction current in isolated olfactory sensory neurons would produce a very low probability of measuring odorant responses (Lagostena & Menini, 2003), due to the fact that each olfactory sensory neuron expresses only one of more than a thousand odorant receptors (for reviews,

see Rodriguez, 2007; Malnic, 2007). To activate CNG channels in the cilia, we included caged 8-Br-cAMP in the intracellular solution filling the patch pipette and applied ultraviolet light flashes to the ciliary region to release the physiologically active 8-Br-cAMP. Upon flash photolysis, CNG channels are activated by 8-Br-cAMP allowing the flux of Ca²⁺ ions in the cilia and the subsequent opening of Ca²⁺-activated Cl⁻ channels (Boccaccio *et al.* 2006; Boccaccio & Menini, 2007). We have previously shown that the rising phase of the response at -50 mV in Ringer solution containing 1 mm Ca²⁺ was multiphasic, composed of a primary phase of the response due to Na⁺ and Ca²⁺ influx through CNG channels and a secondary phase due to Cl⁻ efflux through Cl⁻ channels activated by the influx of Ca²⁺. Moreover, the secondary phase of the response was absent in low extracellular Ca²⁺ or at +50 mV, when the influx of Ca²⁺ through CNG channels is strongly reduced and therefore the contribution of Ca²⁺-activated Cl⁻ channels is expected to be negligible (Boccaccio & Menini, 2007).

To investigate the Ca^{2+} dependence of the rising phase of the response in WT and KO mice, we compared responses at -50 mV in extracellular low Ca^{2+} or in 1 mM Ca^{2+} in the same neuron (Fig. 4*A* and *B*). Both in WT and in KO mice the rising phase of the response in low Ca^{2+} was well

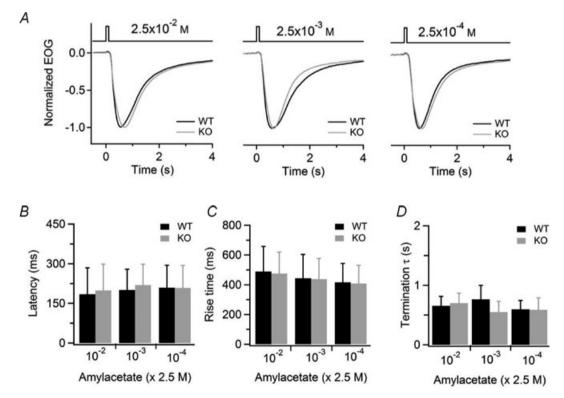


Figure 3. Kinetics analysis of odorant responses in WT and KO mice A, normalized EOG responses to 100 ms pulses of vapour of the indicated amyl acetate concentration in solution for WT (black traces) or KO (grey traces). B–D, average values for latency (B), rise time (C), and time constant of the termination phase (D) were not significantly different in WT and KO animals at each odorant concentration (D = 10).

fitted with a single exponential function, while in Ringer solution containing 1 mm Ca²⁺, the rising phase was slower and could not be described by a single exponential function. To better illustrate the rising phase, traces were normalized to their peak values and plotted superimposed on an expanded time scale in the insets of Fig. 4*A* and *B*. We measured the time necessary for the current to reach 50% of its maximal response, t_{50} , after the delivery of the light flash and found that the average ratio between t_{50} measured in Ringer solution and in low Ca²⁺ at -50 mV was 7.1 ± 3.0 (n = 4) for WT, not significantly different from the value of 9.1 ± 3.2 (n = 5) for KO.

To further investigate the presence of a Ca^{2+} -activated Cl^- current, we compared currents in Ringer solution containing 1 mM Ca^{2+} at +50 or -50 mV in the same

neuron (Fig. 4*C* and *D*). At +50 mV the influx of Ca²⁺ through CNG channels is greatly reduced and the outward current is mainly carried by K⁺ ions, whose permeation through CNG channels is similar to that of Na⁺ ions (reviewed in Kaupp & Seifert, 2002). Both in WT and KO, the rising phase at +50 mV could be well described by a single exponential function, whereas more than one current component was present at -50 mV, as discussed above. The different rising components are illustrated in more detail in the insets of Fig. 4*C* and *D*. The rising time of the response was measured, as described above, as t_{50} , and we found that the ratio between t_{50} at +50 and at -50 mV was 0.27 ± 0.13 (n = 3) for WT, not significantly different from the value of 0.22 ± 0.07 (n = 4) for KO. Furthermore, in the same set of experiments, the ratio

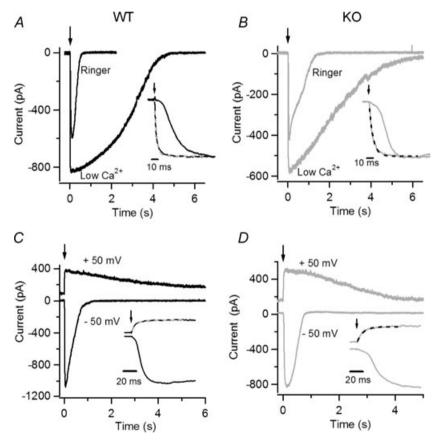


Figure 4. Current responses induced by photorelease of 8-Br-cAMP in isolated olfactory sensory neurons from WT and KO mice

Currents recorded from isolated mouse olfactory sensory neurons in the whole-cell voltage-clamp configuration in response to photorelease of 8-Br-cAMP in the cilia. An ultraviolet flash was applied at the time indicated by each arrow. A and B, an isolated olfactory sensory neuron from WT (A, black traces) and KO (B, grey traces) mice was bathed in Ringer solution containing 1 mm Ca²⁺ or in nominally 0 Ca²⁺ at the holding potential of -50 mV. Insets: responses were scaled to their maximum value and plotted superimposed on an expanded time scale. The rising phase of the response in 0 Ca²⁺ solution was fast and was well fitted by a single exponential (dashed lines), with $\tau = 3.6$ ms for WT and $\tau = 6.2$ ms for KO. Traces in Ringer solution and 0 Ca²⁺ in each panel were recorded from the same neuron. C and D, currents in an isolated olfactory sensory neuron from WT (C, black traces) and KO (D, grey traces) mice at the holding potential of -50 or +50 mV. Insets: current responses plotted on an expanded time scale, displayed a multiphasic rising phase at -50 mV, whereas at +50 mV the rising phase was well fitted by a single exponential (dashed lines, $\tau = 7.7$ ms for WT and $\tau = 10.9$ ms for KO mice). Traces at -50 and +50 mV in each panel were recorded from the same neuron.

between peak current amplitude at +50 and -50 mV was 0.35 ± 0.08 (n = 3) for WT, not significantly different from 0.43 ± 0.30 (n = 4) for KO mice.

These results show that in isolated olfactory sensory neurons from both mouse lines the transduction current comprises a primary CNG current and a secondary Ca²⁺-activated current that is expected to be carried by Cl⁻ ions (Boccaccio & Menini, 2007).

To directly measure Ca^{2+} -activated currents in WT and KO mice, we photoreleased Ca^{2+} in the cilia (Fig. 5). To determine if Ca^{2+} -activated currents were carried by Cl^- , we measured the reversal potentials in the presence of various Cl^- concentrations. In a first set of experiments, we measured the reversal potential in almost symmetrical Cl^- solutions (Fig. 5*A* and *B*), while in a second set of experiments we reduced the intracellular Cl^- concentration by replacing most Cl^- with gluconate (Fig. 5*C* and *D*). The average reversal potential in symmetrical Cl^- solutions for WT, -0.8 ± 1.6 mV (n = 5), was not significantly different from that measured

in KO, -0.5 ± 3.2 mV (n=4) (Fig. 5E). The average reversal potential in the low intracellular Cl⁻ solution was shifted toward more negative values, as expected for Cl⁻ channels in our ionic conditions, and was similar for WT, -37.0 ± 1.2 mV (n=5), and KO, -41.7 ± 5.6 mV (n=5) (Fig. 5E). These results confirm that Ca²⁺-activated Cl⁻ channels were present in olfactory sensory neurons from both WT and KO mice.

Currents in inside-out excised membrane patches from dendritic knob/cilia

To obtain a precise control of the concentrations of cyclic nucleotides and Ca²⁺ at the intracellular side of the transduction channels, we conducted patch-clamp experiments on excised membrane patches from the dendritic knob of dissociated olfactory sensory neurons with visible cilia of WT and KO mice. As previously noted (Reisert *et al.* 2003), cilia were sometimes sucked into the tip of the patch pipette and therefore the excised patches

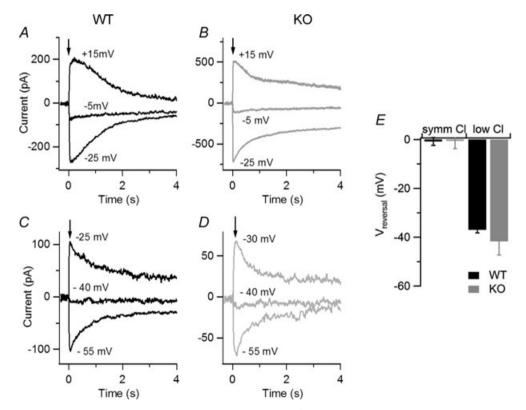


Figure 5. Current responses induced by photorelease of Ca²⁺ in isolated olfactory sensory neurons from WT and KO mice

Currents recorded from isolated mouse olfactory sensory neurons in the whole-cell voltage-clamp configuration in response to photorelease of caged Ca^{2+} (DMNP-EDTA) in the cilia. An ultraviolet flash was applied at the time indicated by each arrow to release the physiologically active Ca^{2+} into the ciliary region. A and B, currents from olfactory sensory neurons were recorded in symmetrical CI^- solutions from WT (A, black traces) and KO (B, grey traces) mice. Currents in each panel were evoked on the same isolated olfactory sensory neuron at the indicated holding potentials, corrected for junction potentials. C and D, similar experiments were repeated when most (see Methods section) CI^- in the intracellular solution was replaced with gluconate. E, average reversal potentials in symmetrical CI^- solutions and in low CI^- solutions.

contained membranes from both the dendritic knob and from the cilia.

To investigate the presence of CNG and Ca²⁺-activated Cl⁻ currents, inside-out excised membrane patches, at

the holding potential of $-50 \,\text{mV}$, were first exposed to a solution containing $100 \,\mu\text{M}$ cAMP in the absence of divalent cations. The same patch was then exposed to a solution containing $100 \,\mu\text{M}$ Ca²⁺ (Fig. 6A and B).

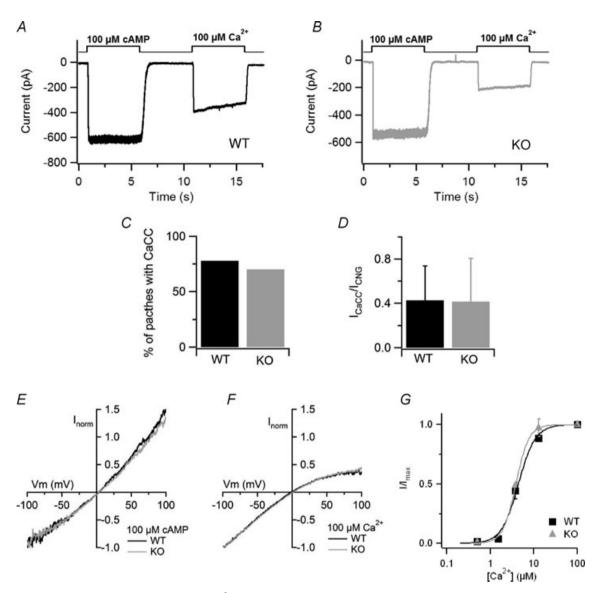


Figure 6. Recordings of CNG and Ca²⁺-activated Cl⁻ currents in inside-out membrane patches

A and B, the cytoplasmic side of membrane patches excised from dendritic knob/cilia of olfactory sensory neurons from WT mice (A) and KO mice (B) was exposed to $100~\mu\text{M}$ cAMP, in the absence of divalent cations, to activate the CNG channels, and to $100~\mu\text{M}$ Ca²⁺ to activate the Cl⁻ channels. Divalent cations were absent from the patch pipette solution. The holding potential was -50~mV. C, percentage of membrane patches with detectable Ca²⁺-activated Cl⁻ currents with respect to the presence of CNG currents in WT and KO mice. D, average ratios between Ca²⁺-activated Cl⁻ currents and CNG currents in patches from WT (n=6) and KO (n=11) mice. E, comparison of representative current–voltage relations of the CNG current activated by $100~\mu\text{M}$ cAMP in WT (black trace) or KO (grey trace) patches. Voltage ramp from -100~to+100~mV. Currents were normalized to the value at -100~mV. F, comparison of representative current–voltage relations of the Cl⁻ current activated by $100~\mu\text{M}$ Ca²⁺ in WT (black trace) or KO (grey trace) patches. Voltage ramps from -100~to+100~mV. Currents were normalized to the value at -100~mV. G, dose–response relations were measured exposing patches to various free Ca²⁺ concentrations. The holding potential was -50~mV. Peak currents at each Ca²⁺ concentrations were normalized to the average current measured in the presence of $100~\mu\text{M}$ Ca²⁺ before and after each test Ca²⁺ concentration. Normalized currents were plotted *versus* Ca²⁺ concentrations, and fitted to the Hill equation. For WT mice $K_{1/2}$ was $4.4~\mu\text{M}$, and n_{H} was 2.2~(n=3). For KO mice $K_{1/2}$ was $3.8~\mu\text{M}$, and n_{H} was 2.9~(n=3).

We found that, both in WT and KO mice, about 75% of the membrane patches that showed detectable CNG currents also had Ca²⁺-activated Cl⁻ currents (Fig. 6C). We observed a great variability in current amplitudes in both mouse lines: at $-50 \,\mathrm{mV}$, currents in WT mice varied from absolute values of 29 to 795 pA for CNG currents, and from 16 to 495 pA for Ca²⁺-activated Cl⁻ currents, while currents in KO mice varied from absolute values of 56 to 902 pA for CNG currents, and from 9 to 212 pA for Ca²⁺-activated Cl⁻ currents. Since it has been previously shown that Ca²⁺-activated Cl⁻ currents in olfactory sensory neurons exhibit a rundown over time, while CNG currents remain quite constant (Reisert et al. 2003), both currents were measured within 30 s after patch excision. The average CNG current at -50 mV was $327 \pm 305 \text{ pA}$ (n=7) in WT animals, not significantly different from the value of $209 \pm 254 \,\mathrm{pA}$ (n=14) in KO mice. For Ca²⁺-activated Cl⁻ currents the average amplitude at $-50 \,\mathrm{mV}$ was $113 \pm 15 \,\mathrm{pA}$ (n=7) in WT animals, also not significantly different from the value of 44 ± 64 pA (n = 14) in KO mice. To obtain an estimate of the relative density of channels we measured the ratio between Ca²⁺-activated Cl⁻ currents and CNG currents in each membrane patch. The average ratio between Ca2+-activated Cl- currents and CNG currents was 0.42 ± 0.31 (n = 7) in WT animals, not significantly different from the value of 0.41 ± 0.39 (n = 14) calculated in KO mice (Fig. 6D). These results show that Ca²⁺-activated Cl⁻ currents are present in inside-out patches from the dendritic knob/cilia of KO mice.

To further compare other biophysical properties of the transduction channels in WT and KO mice, we measured the rectification properties of the two types of currents. Currents were activated by 100 μ M cAMP (Fig. 6E) or by $100 \,\mu\text{M} \,\text{Ca}^{2+}$ (Fig. 6F) using voltage ramps from -100to $+100 \,\mathrm{mV}$ and normalized to the value at $-100 \,\mathrm{mV}$. As shown in Fig. 6E and F, normalized current-voltage relations from WT and KO mice superimposed. The average ratio between currents at +50 and -50 mV for CNG currents was 1.13 ± 0.15 (n = 5) in WT animals, not significantly different from the value of 1.15 ± 0.15 (n=10) calculated in KO mice. For Ca²⁺-activated Cl⁻ currents the average ratio between currents at +50 and -50 mV was 0.52 ± 0.20 (n = 5) in WT animals, also not significantly different from the value of 0.52 ± 0.08 (n = 6) in KO mice.

Since it has been previously shown (Reisert *et al.* 2003) that, in addition to rundown, Ca^{2+} -activated Cl^- currents in olfactory sensory neurons also exhibit a reversible time-dependent decrease in amplitude during the exposure to a constant Ca^{2+} concentration (Fig. 6*A* and *B*), we measured the ratio between current amplitudes measured at the peak and after 5 s of $100 \, \mu M$ Ca^{2+} exposure. We found that the ratio was 0.80 ± 0.10 (n = 6)

in WT mice, not significantly different from the value of 0.78 ± 0.13 in KO animals (n = 12).

Finally, to measure the Ca²⁺ sensitivity of the Cl⁻ channel in WT and KO mice, we obtained dose-response relations by activating currents with various Ca²⁺ concentrations in excised membrane patches (Fig. 6G). Experiments were performed after the rapid phase of rundown, when currents reached an almost steady-state value and, to take into account a remaining slow phase of the rundown that was present in some membrane patches, currents at each test Ca²⁺ concentration were normalized to the average current activated by $100 \,\mu\text{M}$ Ca²⁺ before and after each test Ca²⁺ concentration. Normalized currents measured at -50 mV were plotted versus Ca²⁺ concentration and fitted by the Hill equation: $I/I_{\text{max}} =$ $c^{n_{\rm H}}/(c^{n_{\rm H}}+K_{1/2}^{n_{\rm H}})$, where c is the Ca²⁺ concentration, $K_{1/2}$ the Ca²⁺ concentration producing half-maximal current activation, and $n_{\rm H}$ is the Hill coefficient. At -50 mV, the $K_{1/2}$ was 4.4 μ M for WT, similar to 3.8 μ M for KO mice, and $n_{\rm H}$ was 2.2 and 2.9 for WT and KO animals, respectively (Fig. 6G). $K_{1/2}$ and $n_{\rm H}$ values were not significantly different between WT and KO mice.

Thus, electrophysiological properties of Ca²⁺-activated Cl⁻ currents in inside-out membrane patches from dendritic knob/cilia of olfactory sensory neurons are not significantly different between WT and KO mice.

Discussion

Expression of Best2 in the ciliary layer of the olfactory epithelium

In this study, we confirmed that Best2 is expressed in the ciliary layer of olfactory sensory neurons and that the antibody against Best2 we previously developed (Pifferi et al. 2006b) is specific for this protein, as demonstrated by the absence of immunostaining in the olfactory epithelium of KO mice. Thus, these data, together with those of a very recently published study (Klimmeck et al. 2009) in which the expression of Best2 in the olfactory epithelium was extensively investigated, solve the reported controversy (Hartzell et al. 2008; Marmorstein et al. 2009) as to whether mouseBest2 is indeed expressed in the olfactory epithelium. In fact, Bakall et al. (2008), by employing immunoistochemistry in the same mouse lines of the present study, found expression of the Best2 protein in the colon and in the eye (in the non-pigmented epithelia cells of the ciliary body) of WT mice, but not in the olfactory epithelium, although they detected a Best2 transcript. A possible explanation of the different results between the laboratories is likely to reside in the different antibodies that were used. Indeed, Bakall et al. (2008) reported that their antibody was not working in Western blot. On the other side, concerning the immunohistochemistry, it must be noted that no positive controls of the immunostaining in the olfactory epithelium were shown by Bakall *et al.* (2008) and therefore it cannot be excluded that, since cilia are very fragile, they were absent from the olfactory epithelium slices.

Physiological role of Best2

Since Best2 is expressed in the cilia of olfactory sensory neurons (Pifferi et al. 2006b), the site of olfactory transduction, and it has been previously demonstrated that Best2 forms Ca²⁺-activated Cl⁻ channels when expressed in heterologous systems (Qu et al. 2004; Qu & Hartzell, 2004; Pifferi et al. 2006b), we conducted experiments to investigate the physiological role of Best2 in olfactory transduction by comparing the properties of WT mice with those of KO mice. Since a previous study has shown that KO mice showed no obvious olfactory deficits, as detected by the cookie test (Bakall et al. 2008), we also repeated this type of behavioural test and found results in agreement with those previously published. However, it cannot be concluded from these experiments that Best2 is not required for normal olfactory sensitivity, as more detailed behavioural studies, as for example the study of olfactory behavioural thresholds as reported in mice lacking NKCC1 (Smith et al. 2008), would be required to reach such a conclusion. In addition, it is important to note that, as previously pointed out by Smith et al. (2008), it is still unknown whether a deficit in Ca2+-activated Cl⁻ current would produce a reduction in olfactory sensitivity, and it is possible that the secondary Cl⁻ current may not be required at all for normal olfactory sensitivity. Indeed, it is possible that the primary current through CNG channels is sufficient for normal olfactory sensitivity.

We analysed the responses to odorants of the olfactory epithelium and found no differences in the odorant sensitivity or kinetics properties measured by EOG recordings in WT and KO mice. Moreover, at the level of single olfactory sensory neurons, a Ca²⁺-activated Cl⁻ channel component was also measured by using photolysis of caged 8-Br-cAMP or of caged Ca²⁺ localized to the cilia of olfactory sensory neurons of KO mice. Finally, a Ca²⁺-activated Cl⁻ current was still present in excised inside-out patches from knob/cilia of olfactory sensory neurons of KO mice, with electrophysiological properties similar to those of WT mice.

Thus, we determined that the absence of expression of Best2 in the ciliary layer of the olfactory epithelium does not significantly alter the electrophysiological properties of the olfactory epithelium. These results indicate that Best2 may not be the main molecular component of the olfactory Ca²⁺-activated Cl⁻ channel, although we cannot exclude the possibility that in KO mice some compensatory mechanisms may act to replace the function of the missing protein. However, it must also

be noted here that, although the protein Best2 has been proposed as a candidate for being a molecular component of the olfactory Ca²⁺-activated Cl⁻ channel, very recent studies reported that the anoctamin/TMEM16 family of membrane proteins display many features of native Ca²⁺-activated Cl⁻ channels (Caputo et al. 2008; Schroeder et al. 2008; Yang et al. 2008). In addition, it has been shown, by in situ hybridization, that TMEM16B is expressed in the mature sensory neurons of the mouse olfactory epithelium (Yu et al. 2005), and it is a prominent protein in the rat olfactory ciliary proteome (transmembrane protein 16B isoform 2, Supplementary Table S1 in Mayer et al. 2009). Stephan et al. (2009) also identified TMEM16B (Anoctamin 2, ANO2) in a proteomic screen of ciliary membranes and showed that the fusion protein TMEM16B-EGFP localized to the cilia when expressed in vivo using an adenoviral vector. In addition, Stephan et al. (2009) provided evidence that the electrophysiological properties of this protein are remarkably similar to those of native olfactory Ca²⁺-activated Cl⁻ channels.

We have also recently characterized the electrophysiological properties of the mouse TMEM16B expressed in a heterologous system and found that the channel properties are remarkably similar to those of the native Ca²⁺-activated Cl⁻ channels (Pifferi et al. 2009). Indeed, while we have previously pointed out that a significant difference between Best2 and the native olfactory channel was a Ca2+ sensitivity difference of one order of magnitude, with a Ca²⁺ concentration for half-maximal activation at $-50 \,\mathrm{mV}$ of $0.4 \,\mu\mathrm{M}$ for Best2 and a higher concentration of 4.7 μ M for native channels (Pifferi et al. 2006b), we recently showed that the $K_{1/2}$ for Ca²⁺ in TMEM16B-induced currents was 4.9 μ M, similar to the native channel. Further studies will have to establish whether TMEM16B is a component of the olfactory Ca²⁺-activated Cl⁻ channel.

In this study, we have shown that Best2 is expressed in the olfactory ciliary layer, but it does not appear to be the main molecular component of the Ca²⁺-activated Cl⁻ current. What is then the physiological role of Best2 in the olfactory epithelium? In addition to functioning as Ca²⁺-activated Cl⁻ channels when expressed heterologously, mouse Best2 and other bestrophins have also been shown to be activated by osmotic cell swelling in the absence of Ca²⁺, indicating that they may be cell volume regulators (Fischmeister & Hartzell, 2005; Chien & Hartzell, 2007). Furthermore, Best1 can interact with the voltage-gated Ca²⁺ channel Cav1.3 and modulate its biophysical properties (Rosenthal et al. 2006; Yu et al. 2008), indicating that some bestrophins could also act as regulators of other ion channels. In the eye, Best2 is expressed in the non-pigmented epithelium and, using the same mouse line employed in the present study, it was found that KO mice have a diminished

intraocular pressure compared to WT mice (Bakall *et al.* 2008). A very recent study suggested that Best2 is involved in the control of aqueous dynamics (Zhang *et al.* 2009).

In the olfactory epithelium, Klimmeck et al. (2009) have recently investigated in detail the expression pattern of Best2 comparing results from adult and postnatal day 1 (P1) mice. In adult mice, whose olfactory epithelium contains mainly mature olfactory sensory neurons, Best2 expression was found in the cilia of these neurons. In P1 mice, where most olfactory sensory neurons are still immature, strong Best2 signals were detected at all subcellular levels of the developing neurons. Based on these observations, Klimmeck et al. (2009) suggested a physiological role for Best2 related to neurogenesis, in which Best2 may act as a volume-regulated anion channel contributing to the coordinated extension of cell volume in developing sensory neurons.

In the olfactory cilia, it is possible to speculate that Best2 may subserve different functions as, for example, the maintenance of the local ionic homeostasis, also preventing volume changes resulting from the exposure to exogenous osmotically active solutions. It is also possible to envisage that Best2 could contribute to setting the correct chloride concentration in the mucous layer of the olfactory epithelium.

In summary, we now have unequivocal confirmation that Best2 is expressed in the cilia of mature olfactory sensory neurons, thus cutting out all the controversies about its presence in the olfactory epithelium. Nevertheless, the function of Best2 remains elusive and further studies will be required to determine its physiological role.

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Author contributions

S.P., M.D., A.B., A.AQ., R.T. and A.M. contributed to conception and design of the experiments, drafting and revision of the manuscript. M.D. did the immunohistochemistry. F.G. and R.T. did the behavioural test. A.AQ performed RT-PCR and WB analysis. S.P., C.S. and A.B. did the electrophysiology experiments.

S.P., C.S., A.B. and A.M. analyzed the data. S.P., M.D., R.T., A.B. and A.M. wrote the manuscript. All authors approved the final version to be published.

Acknowledgements

We thank C. Degrassi and M. Stebel for animal care; J. Franzot for genotyping mice; L. Masten for technical help; F. Müller and U. B. Kaupp (Forschungszentrum Jülich, Jülich, Germany) for the gift of the CNGA2 monoclonal antibody; V. Hagen (Leibniz-Institut für Molekulare Pharmakologie, Berlin, Germany) for kindly

providing caged cyclic nucleotides. This study was supported by grants from the Italian Ministry of Research (MIUR) and from the Italian Institute of Technology.

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3.2 Calcium concentration jumps reveal dynamic ion selectivity of calcium-activated chloride currents in mouse olfactory sensory neurons and TMEM16b-transfected HEK 293T cells.

Sagheddu C, Boccaccio A, Dibattista M, Montani G, Tirindelli R, Menini A (2010). J Physiol. 588, 4189–4204

Calcium concentration jumps reveal dynamic ion selectivity of calcium-activated chloride currents in mouse olfactory sensory neurons and TMEM16b-transfected HEK 293T cells

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Ca²⁺-activated Cl⁻ channels play relevant roles in several physiological processes, including olfactory transduction, but their molecular identity is still unclear. Recent evidence suggests that members of the transmembrane 16 (TMEM16, also named anoctamin) family form Ca²⁺-activated Cl⁻ channels in several cell types. In vertebrate olfactory transduction, TMEM16b/anoctamin2 has been proposed as the major molecular component of Ca²⁺-activated Cl⁻ channels. However, a comparison of the functional properties in the whole-cell configuration between the native and the candidate channel has not yet been performed. In this study, we have used the whole-cell voltage-clamp technique to measure functional properties of the native channel in mouse isolated olfactory sensory neurons and compare them with those of mouse TMEM16b/anoctamin2 expressed in HEK 293T cells. We directly activated channels by rapid and reproducible intracellular Ca²⁺ concentration jumps obtained from photorelease of caged Ca²⁺ and determined extracellular blocking properties and anion selectivity of the channels. We found that the Cl⁻ channel blockers niflumic acid, 5-nitro-2-(3-phenylpropylamino) benzoic acid (NPPB) and DIDS applied at the extracellular side of the membrane caused a similar inhibition of the two currents. Anion selectivity measured exchanging external ions and revealed that, in both types of currents, the reversal potential for some anions was time dependent. Furthermore, we confirmed by immunohistochemistry that TMEM16b/anoctamin2 largely co-localized with adenylyl cyclase III at the surface of the olfactory epithelium. Therefore, we conclude that the measured electrophysiological properties in the whole-cell configuration are largely similar, and further indicate that TMEM16b/anoctamin2 is likely to be a major subunit of the native olfactory Ca²⁺-activated Cl⁻ current.

(Received 17 June 2010; accepted after revision 8 September 2010; first published online 13 September 2010)

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Abbreviations DIDS, 4,4'-diisothiocyanatostilbene-2,2'-disulfonic acid; HEK, human embryonic kidney; MeS⁻, methanesulfonate; NFA, niflumic acid; NPPB, 5-nitro-2-(3-phenylpropylamino)benzoic acid; SCN⁻, isothiocyanate; TMEM16, transmembrane 16.

Introduction

In several cell types, an increase in intracellular Ca²⁺ concentration produces the activation of chloride channels that, depending on the electrochemical gradient of Cl⁻, will cause depolarization or hyperpolarization

of the cell membrane. Ca²⁺-activated Cl⁻ channels were first identified in *Xenopus* oocytes (Miledi, 1982; Barish, 1983) and in the inner segment of salamander photoreceptors (Bader *et al.* 1982), and afterwards in many other cell types, including olfactory sensory neurons (Kleene & Gesteland, 1991; Kleene, 1993; Kurahashi

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& Yau, 1993). These channels are involved in a large variety of physiological processes, including generation of the fertilization potential in *Xenopus* oocytes, regulation of synaptic transmission in photoreceptors, and signal amplification in olfactory sensory neurons (reviewed by Frings *et al.* 2000; Hartzell *et al.* 2005; Kleene, 2008; Frings, 2009*a*,*b*).

Despite the physiological relevance of Ca²⁺-activated Cl⁻ channels, their molecular identity remained largely elusive. Several molecular candidates have been proposed for these channels, but none of them completely reproduced the properties of native Ca²⁺-activated Cl⁻ currents (reviewed by Hartzell *et al.* 2005; Duran *et al.* 2010). In 2008, three independent studies reported evidence suggesting that some members of the family of TMEM16/anoctamins are likely to be the molecular determinants of Ca²⁺-activated Cl⁻ currents in some cell types (Caputo *et al.* 2008; Schroeder *et al.* 2008; Yang *et al.* 2008; reviewed by Flores *et al.* 2009; Galietta, 2009; Hartzell *et al.* 2009; Kunzelmann *et al.* 2009).

In olfactory sensory neurons, Ca²⁺-activated Cl⁻ currents are measured, together with cAMP-activated currents, in the cilia (Kleene & Gesteland, 1991; Kleene, 1993), where they play an important role in the amplification of the response to odorants, constituting up to 90% of the transduction current (Kurahashi & Yau, 1993; Lowe & Gold, 1993; Boccaccio & Menini, 2007). Indeed, the process of olfactory transduction occurs in the cilia of olfactory sensory neurons, where a second messenger cascade is activated by the binding of odorant molecules to odorant receptors and leads to the production of cAMP and the opening of cAMP-activated channels (reviewed by Schild & Restrepo, 1998; Lowe & Gold, 1993; Menini, 1999; Matthews & Reisert, 2003; Menini et al. 2004; Pifferi et al. 2006a, 2009c; Kleene, 2008; Tirindelli et al. 2009). Since olfactory sensory neurons maintain an unusually elevated intracellular concentration of Cl-(Reuter et al. 1998; Kaneko et al. 2001, 2004), the influx of Ca²⁺ through cAMP-activated channels in the cilia produces an efflux of Cl- through Ca²⁺-activated Cl- channels, contributing to the odorant-induced depolarization (Kleene & Gesteland, 1991; Kleene, 1993, 1997, 2008; Kurahashi & Yau, 1993; Lowe & Gold, 1993; Boccaccio & Menini, 2007; reviewed by Frings et al. 2000; Frings, 2009*a*,*b*; Pifferi *et al.* 2009*c*).

At present, several lines of evidence indicate that TMEM16b/anoctamin2 is the best candidate for being the main molecular component of the olfactory Ca²⁺-activated Cl⁻ channels in the cilia. Indeed, *in situ* hybridization studies showed that TMEM16b/anoctamin2 is expressed in mature sensory neurons of the mouse olfactory epithelium (Yu *et al.* 2005); proteomic screenings identified TMEM16b/anoctamin2 as a prominent protein of olfactory ciliary membranes (Stephan *et al.* 2009;

Hengl *et al.* 2010; Rasche *et al.* 2010); the fusion protein TMEM16b/anoctamin2–EGFP localized to the cilia when expressed *in vivo* using an adenoviral vector (Stephan *et al.* 2009); immunohistochemistry showed the localization of TMEM16b/anoctamin2 to the ciliary region (Hengl *et al.* 2010; Rasche *et al.* 2010); functional properties measured by patch-clamp recordings from excised inside-out membrane patches of TMEM16b/anoctamin2 expressed in HEK 293T cells or from the dendritic knobs and ciliary region of olfactory sensory neurons are very similar (Pifferi *et al.* 2009*a*; Stephan *et al.* 2009).

However, to identify the channel protein it is necessary to prove that all the functional properties of native channels are reproduced by the candidate protein. At present, several electrophysiological properties of native olfactory Ca²⁺-activated Cl⁻ currents are still unknown. Indeed, while the properties of native olfactory channels in the excised cilium (Kleene & Gesteland, 1991; Kleene, 1993) or in the excised inside-out membrane patches have been extensively investigated (Reisert et al. 2003; Pifferi et al. 2006b, 2009b; Stephan et al. 2009), those of the native channels in isolated olfactory sensory neurons are poorly known. Moreover, currents in excised patches exhibited a pronounced rundown as well as inactivation/desensitization in the presence of a constant Ca²⁺ concentration (Reisert et al. 2003), while whole-cell recordings appeared to be more stable (Boccaccio & Menini, 2007; Takeuchi et al. 2009).

Niflumic acid or 4-acetamido-4-isothiocyanato-stilben-2, 2-disulfonate (SITS; Kurahashi & Yau, 1993; Lowe and Gold, 1993) are commonly used as extracellular blockers of Ca²⁺-activated Cl⁻ channels in intact olfactory sensory neurons, but the extracellular blocking potencies of several other compounds have not been measured. Moreover, the ion selectivity of the native channels in isolated olfactory sensory neurons has not been estimated yet, except for showing that the current is carried by Cl⁻ ions (Kurahashi & Yau, 1993; Takeuchi *et al.* 2009).

The goal of this study was to measure the unknown electrophysiological properties of the native olfactory Ca²⁺-activated Cl⁻ channels and to obtain a side-by-side comparison with the recently cloned TMEM16b/anoctamin2 (Stephan *et al.* 2009) heterologously expressed in HEK 293T cells. Channels were directly activated by rapidly increasing the intracellular Ca²⁺ concentration by flash photolysis of caged Ca²⁺.

Our results show that Ca²⁺-activated Cl⁻ currents measured in the whole-cell configuration in olfactory sensory neurons were largely similar to those induced by the candidate protein, contributing further support to the hypothesis that TMEM16b/anoctamin2 is a major constituent of the olfactory Ca²⁺-activated Cl⁻ channel.

Methods

Ethical approval

All animals were handled in accordance with the Italian Guidelines for the Use of Laboratory Animals (Decreto Legislativo 27/01/1992, no. 116) and European Union guidelines on animal research (No. 86/609/EEC). For experiments mice were killed by cervical dislocation or anaesthetized with CO₂ inhalation and then decapitated.

RNA extraction and RT-PCR

RNA was extracted from the olfactory epithelium of FVB mice and purified using Trizol reagent (Invitrogen Milano, Italy). About 2 μ g of total RNA served as template for oligo-dT primed first strand cDNA synthesis with Im-Prom-II Reverse Transcriptase (Promega, Milano, Italy). PCR was performed in Mastercycler Personal (Eppendorf, Milano, Italy) using AmpliBiotherm DNA polymerase, 3 mm MgCl₂, 0.2 mm for each dNTPs and 200 pmol forward/reverse target-specific oligonucleotide primers. Cycling parameters consisted of an initial denaturation step (95°C, 2 min) followed by 35 cycles each of these included a denaturation (95°C, 30 s), a primer annealing (50° C, 30 s), and an extension (72° C, 30 s) step. Reaction was completed by a final extension step at 72°C for 5 min. Semiquantitative analysis of RNA expression was performed on agarose gel after electrophoresis using the NIS-Elements Advanced Research software (Nikon, Firenze, Italy).

Primers were designed to amplify a 650–700 bp DNA sequence, which is predicted to encode a region of the C-terminal intracellular domain of TMEM16/anoctamins. The chromosomic region corresponding to the amplicon DNA sequences spans over five exons and four introns.

The following primer sequences were used to amplify target DNAs:

TMEM16a/anoctamin1: 1826-2493, fwd: bases 5'-ACGTGTACATCTTCCGCTCTTT-3' (Tm 58°C), rev: 5' GATCTGAACCTCATAGCCCAG-3' (Tm 59°C); TMEM16b/anoctamin2: 1694-2358, bases fwd: 5'-ATGTCTACGTGTTCGACGGTTA-3' (Tm 58°C) rev: 5'-AAACTGAACCTCCTGGTCGAA (Tm 57°C); TMEM16c/anoctamin3: bases 2009-2652, fwd: 5'-ACAATAAACTTTTTGAGCGGTG-3' (Tm 54°C) rev: 5'-GTAACCAGATTTTCCCATACC-3' (Tm 55°C); TMEM16d/anoctamin4: bases 1361-2052, fwd: 5'-ACTTGAGATTGATAAACAGGTG-3' (Tm 54°C) rev: 5'-GTACTTCAGAGGGGTTCCTGA-3' (Tm 59°C); TMEM16e/anoctamin5: bases 1739–2382, fwd: 5'-ACACATATGTTCAACATATGGA-3' (Tm 54°C), rev: 5'-GGTGACGAAGTCTTTTTTCTC-3') (Tm 55°C);

TMEM16f/anoctamin6: bases 1751-2418, fwd: 5'-CAGTGTACTTGCTGGGCAAATA-3' (Tm 57°C), rev: 5'-CAAGGTATAGTTACCAAGCCC-3'(Tm 57°C); TMEM16g/anoctamin7: bases 1685-2340, fwd: 5'-ACCACACCTTGTTTGGAATCC-3' (Tm 57°C), rev: 5'-GTAAGTCGGAGAATAGTGTCC 3' (Tm 57°C); TMEM16h/anoctamin8: bases 1962-2665, fwd: 5'-CGAAGAAGACGATGAGCCTGA-3' (Tm 71°C), rev: 5'-CCTGCCGCTCGTGCCGCTTGA-3' (Tm 61°C); TMEM16j/anoctamin9: bases 1331-2004, fwd: 5'-CCACGCGCCTGGCTGGCCTGTG-3' (Tm 63°C), rev: 5'-CACGGTCACGTTTTCTTGGCC-3' (Tm 54°C); TMEM16k/anoctamin10: bases 1298-1894, fwd: 5'-TGGCCACACTCCTGATCACCTC-3' (Tm 59°C), rev: 5'-AAGATTCGAATTCCAATCTGG-3' (Tm 67°C); OMP (olfactory marker protein): bases 1-1079, fwd: 5'-CCCTGCTGGCCAAAGCTGGAA-3' (Tm 63°C), rev: 5'-GTCTCTAAAGCTGTAGGGAGA-3' (Tm 58°C); Vmn2r70, bases 1011-1704, fwd: 5'-TTACAGTAGTG-AATTTTCCTTTGC-3' (Tm 55°C), rev: 5'-TTGGAGGCAGAGAGTATGGTGTTC (Tm 63°C).

All amplicons were separated by agarose gel electrophoresis, the corresponding bands were excised and the DNA extracted and purified (Qiagen gel extraction Kit, Milano, Italy) and subsequently subcloned in pGEMt-easy vector (Promega) for sequencing.

Immunohistochemistry

Immunostainings of olfactory epithelium were performed as previously described (Pifferi *et al.* 2009*b*). Primary antibodies were: mouse monoclonal anti-TMEM16b (1:1; provided by H. Stöhr, Universität Regensburg, Regensburg, Germany; Stöhr *et al.* 2009) and rabbit anti-adenylyl cyclase III (1:300, Santa Cruz Biotechnology, SantaCruz, CA, USA; cat. no. sc-558). Secondary antibodies were: Alexa 488-conjugated goat anti-mouse and Alexa 594-conjugated goat anti-rabbit diluted to 1:300 (Molecular Probes-Invitrogen, West Eugene, OR, USA).

Images were visualized by Leica TCS SP2 confocal microscope (Leica Microsystems, Milano, Italy) acquired using Leica software at 1024×1024 pixels resolution and analysed with ImageJ software. Images were not modified other than to balance brightness and contrast.

Dissociation of mouse olfactory sensory neurons

Olfactory sensory neurons were dissociated enzymatically from the olfactory epithelium of 1- to 3-month-old C57 Black mice with a method similar to that previously described (Lagostena & Menini, 2003; Boccaccio *et al.*

2006). The olfactory epithelium was removed and transferred in 1 ml of zero-divalent mammalian Ringer solution with 200 μ M cystein and 2 U ml⁻¹ papain (Sigma, Milano, Italy) for 8–10 min at room temperature. The olfactory epithelium was minced with fine forceps. The reaction was stopped by adding 0.5 ml of Ringer solution with 0.1 mg ml⁻¹ bovine serum albumin (BSA), 0.3 mg ml⁻¹ leupeptin and 0.02 mg ml⁻¹ of DNaseI (all from Sigma). After centrifugation (300 g for 5 min) the cells were resuspended in 1 ml of Ringer solution and plated on glass coverslips (WPI, Sarasota, FL, USA), coated with poly-L-lysine and concanavalin-A Type V (Sigma). Before use, dissociated olfactory sensory neurons were allowed to settle for 60 min at $+4^{\circ}$ C.

Only olfactory sensory neurons with clearly visible cilia were used for the experiments.

Heterologous expression of TMEM16b/anoctamin2

The full-length, dominant olfactory isoform of the mouse TMEM16b/anoctamin2 cloned into the expression vector pAdtrack-CMV (Stratagene, LaJolla, CA, USA) with an independent expression cassette for EGFP (provided by Haiqing Zhao of the Johns Hopkins University in Baltimore (Stephan *et al.* 2009), was transfected into HEK 293T cells using FuGENE 6 reagent (Roche Applied Science, Mannheim, Germany) according to the manufacturer's protocol. Transfected cells were identified by EGFP fluorescence and used for electrophysiological recordings from 24 to 48 h after transfection.

Patch-clamp recordings

Olfactory sensory neurons or HEK 293T cells transfected with TMEM16b/anoctamin2 were observed with an inverted microscope (Olympus IX70, Milano, Italy) with an oil immersion $\times 100$ objective (Zeiss, Milano, Italy).

Currents in the whole-cell voltage-clamp mode were recorded with an Axopatch 200B patch-clamp amplifier, controlled by Clampex 8 via a Digidata 1322A (Axon Instruments, Union City, CA, USA). Patch pipettes were made using borosilicate capillaries (WPI) and pulled with a Narishige PP83 puller (Narishige, Tokyo, Japan). Pipette resistances were 3–7 M Ω when filled with the standard intracellular solution. Currents were low-pass filtered at 1 kHz and acquired at 2 kHz for experiments with olfactory sensory neurons, or filtered at 5 kHz and sampled at 10 kHz for experiments with transfected HEK 293T cells. All experiments were carried out at room temperature (20–24°C).

Ionic solutions and perfusion system

The extracellular mammalian Ringer solution contained (in mm): 140 NaCl, 5 KCl, 1 CaCl₂, 1 MgCl₂, 10 Hepes, 10 glucose and 1 sodium pyruvate (pH 7.4). The pipette solution contained (in mm): 140 CsCl, 3 DMNP–EDTA, 1.5 CaCl₂ and 10 Hepes (pH 7.4). The caged Ca²⁺ compound DMNP–EDTA was purchased from Molecular Probes–Invitrogen, and CaCl₂ was adjusted with a 0.1 M standard solution from Fluka (Deisenhofen, Germany). Pipette solution aliquots were stored for a few days at –20°C and kept refrigerated in the dark during the experimental session. Caged Ca²⁺ was allowed to diffuse from the patch pipette into the cell cytoplasm for at least 2 min after establishment of the whole-cell configuration.

5-nitro-2-(3-Niflumic acid (NFA) and phenylpropylamino)benzoic acid (NPPB, **Tocris** Bioscience, Bristol, UK) were prepared in dimethyl sulfoxide (DMSO) as stock solutions at 200 mm or 83 mM, respectively, and diluted to the final concentration of $400 \,\mu\text{M}$ and $100 \,\mu\text{M}$, respectively, in the bathing solution (DMSO alone did not alter the currents); 4,4'-diisothiocyanatostilbene-2,2'-disulfonic acid (DIDS) was directly dissolved in the bathing solution to 1 mm.

Different bathing solutions were delivered by using a gravity-fed perfusion system. A slow flow rate was selected in such a way that the position of the cilia of the neurons was not perturbed. A complete solution change was obtained in about 10 s. To measure blocker effects, current recordings were obtained before blocker application (control), 1–2 min after delivery of the solution with the blocker, and 2–5 min after perfusion with Ringer solution without the blocker (washout).

For ionic selectivity experiments, Cl⁻ was substituted with other anions by replacing NaCl on an equimolar basis with NaX, where X is the substituted anion, or NaCl was replaced with equimolar choline chloride.

The bath was grounded through a 1 M KCl agar bridge connected to a Ag-AgCl reference electrode. Liquid junction potentials were calculated using Clampex's Junction Potential Calculator, based on the JPCalc program developed by Barry (1994; see also http://web. med.unsw.edu.au/phbsoft/LJP_Calculator.htm). Applied membrane potentials were corrected off-line. The liquid junction potential between the pipette and the Ringer solution was calculated. Then, if the bathing solution was changed after reaching the whole-cell configuration, we calculated the additional liquid junction potential generated between the bathing solution and the 1 M KCl agar bridge. We corrected membrane potentials for the following calculated liquid junction potentials (in mV) in the indicated bathing solutions: -4.6 in Ringer solution, -4.0 in isothiocyanate Ringer solution, -4.7 in bromide Ringer solution, -4.6 in iodide Ringer solution, -4.3 in nitrate Ringer solution, -3.0 in methanesulfonate Ringer solution, -5.3 in Ringer solution with NaCl replaced with choline chloride.

Chemicals, unless otherwise stated, were purchased from Sigma.

Photolysis of caged Ca²⁺

For flash photolysis of caged Ca^{2+} , we used a xenon flash-lamp JML-C2 system (Rapp OptoElectronic, Hamburg, Germany) coupled with the epifluorescence port of the microscope with a quartz light guide as previously described (Boccaccio *et al.* 2006; Boccaccio & Menini, 2007). The spot of light had a diameter of about 15 μ m and was focused on the ciliary region of olfactory sensory neurons or to cover about 50% of the surface of HEK 293T cells. The flash duration was less than 1.5 ms and was kept constant during each experiment. At the beginning of each experiment, the stability of the response was checked by applying repetitive flashes at intervals of about 2 min. Neurons or cells that did not reach a stable response to two or three flashes were discarded.

Data analysis

Data analysis and figures were made with Igor software (Wavemetrics, Lake Oswego, OR, USA). A single exponential function was fitted to the rising phase for monophasic current responses. Current recordings at each holding potential were plotted by subtracting the value of the baseline.

Data are reported as mean \pm s.E.M. and the total number of cells (n). Statistical significance was determined using unpaired t tests, or ANOVA, as appropriate. When a statistically significant difference was determined with ANOVA, a Tukey *post hoc* test was done to evaluate which data groups showed significant differences. P values < 0.05 were considered significant.

Results

Expression of TMEM16s/anoctamins in the olfactory epithelium

To analyse the expression of each TMEM16/anoctamin in the olfactory epithelium, we performed RT-PCR using intron-spanning primers for amplifying DNA encoding a fragment of the C-terminal intracellular domain of each family member (Fig. 1).

We found expression in the olfactory epithelium for TMEM16a/anoctamin1, b/2, f/6, j/9 and k/10 (Fig. 1A). To semiquantify the expression of the different TMEM16/anoctamin isotypes in the olfactory epithelium, we amplified the OMP sequence in the same reaction.

Results indicated that TMEM16a/anoctamin1, b/2 and j/9 are abundantly expressed, and f/6 and k/10 are moderately expressed (Fig. 1D). To further investigate the expression of TMEM16c/anoctamin3, d/4, e/5, g/7 and h/8 (undetectable in Fig. 1A and D) we loaded them in a much larger amounts and found a positive expression of d/4 and g/7 (Fig. 1B). No expression could be detected for TMEM16c/anoctamin3, e/5 and h/8 (Fig. 1A and B) although they are abundantly expressed in the whole E16 embryo (Fig. 1C). To validate the PCR results, all amplicons were directly sequenced after purification from agarose gel.

To examine the localization of TMEM16b/anoctamin2 in the olfactory epithelium, we performed immunohistochemistry experiments on cryosections of the olfactory epithelium by using an anti-TMEM16b antibody (Stöhr *et al.* 2009). We found staining at the surface of

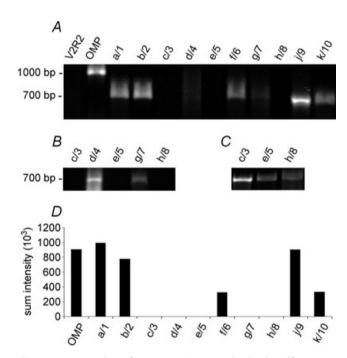


Figure 1. Expression of TMEM16s/anoctamins in the olfactory epithelium

A, olfactory transcripts of the TMEM16/anoctamin isoforms 1–10 were detected by RT-PCR (35 cycles) and sequencing. Specificity was confirmed by negative (olfactory epithelium cDNA amplified with primer for vomeronasal receptors, V2R2s) and positive (olfactory epithelium cDNA amplified with primers for OMP) controls. All lanes were loaded with the same amount of reaction product (5 μ l) to obtain a semiquantitative evaluation of the TMEM16/anoctamin expression as represented in D. B, TMEM16c/anoctamin3, d/4, e/5, g/7 and h/8 (apparently undetectable in A) were loaded in a much larger amount and showed a lower but positive expression of d/4 and g/7. C, primer specificity for TMEM16c/anoctamin3, e/5 and h/8 (not detected in the olfactory epithelium) was confirmed by RT-PCR on cDNA from the whole embryo (E16). D, semiquantitative analysis of the TMEM16/anoctamin expression in the olfactory epithelium for the experiment shown in A.

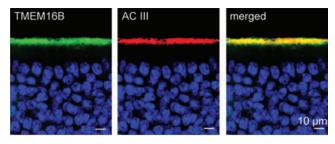


Figure 2. Localization of TMEM16b/anoctamin2 at the surface of the olfactory epithelium

Immunostaining of sections of the olfactory epithelium. Confocal micrographs showing TMEM16b and adenylyl cyclase III (AC3) expression at the surface of the olfactory epithelium. Cell nuclei were stained by DAPI. The image on the right was obtained from the merge of the left and centre images. Scale bar is 10 μ m in all panels.

the olfactory epithelium, where TMEM16b/anoctamin2 largely co-localized with adenylyl cyclase III (Fig. 2), a protein that is mainly expressed in the cilia of olfactory sensory neurons (Menco *et al.* 1992, 1994; reviewed by Menco, 1997). However, we cannot exclude the presence of TMEM16b/anoctamin2 also in the microvilli of sustentacular cells at the surface of the olfactory epithelium. Therefore, our experiments are in agreement with previous results, obtained with different antibodies, showing that TMEM16b/anoctamin2 is expressed at the site of olfactory transduction (Hengl *et al.* 2010; Rasche *et al.* 2010).

Extracellular blockers of native Ca²⁺-activated currents in olfactory sensory neurons The most commonly used extracellular blockers.

The most commonly used extracellular blocker of Ca^{2+} -activated Cl^- current in intact olfactory sensory neurons is NFA at concentrations ranging between 300 and 500 μ M (Kleene, 1993; Boccaccio *et al.* 2006; Boccaccio & Menini, 2007; Takeuchi *et al.* 2009), while the extracellular blocking potencies of several other compounds are still unknown. We measured the effect of adding 400 μ M NFA, 100 μ M NPPB or 1 mM DIDS at the extracellular side of olfactory sensory neurons, while activating the current by producing a sudden Ca^{2+} concentration increase by photorelease of caged Ca^{2+} in the ciliary region at the holding potential of -50 mV.

Figure 3A illustrates the typical NFA blockage of the current activated by a Ca²⁺ concentration jump produced by an ultraviolet flash applied at the time indicated in the upper panels. The maximal current amplitude was reduced to about 17% of its value before blocker application, and the effect was reversed after perfusion with Ringer solution without NFA (87% recovery), in agreement with previous data (Kleene, 1993; Boccaccio et al. 2006; Boccaccio & Menini, 2007; Takeuchi et al. 2009). To better illustrate the variability of results among neurons, we normalized the responses for each neuron in the presence and after washout of the blocker to the value measured before blocker application (Fig. 3, right panels). On average, the amplitude in the presence of 400 μ M NFA was 13% of the control value. After perfusion with Ringer solution without the blocker, the current recovered on average to 63% of its control value.

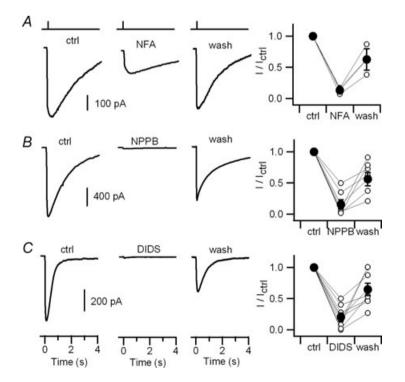


Figure 3. Olfactory sensory neurons: extracellular blockage of currents activated by photorelease of caged Ca²⁺ in the cilia

Currents recorded in the whole-cell voltage-clamp configuration in response to sudden jumps in Ca^{2+} concentration obtained with photorelease of caged Ca^{2+} . Ultraviolet light flashes were applied on the ciliary region at the times indicated by the vertical bars in the upper panels. The holding potential was -50 mV. The following blockers were used: A, $400~\mu$ m NFA; B, $100~\mu$ m NPPB; C, 1~mm DIDS. Current recordings were obtained before blocker application (control), 1-2~min after application of the indicated blockers, and 2-5~min after the removal of blockers (washout). Panels on the right show peak currents measured in the presence of each blocker and after washout, normalized to the control value before blocker application for each neuron (open circles). Average ratios are plotted as filled circles.

Figure 3*B* shows recordings from an olfactory sensory neuron in which the extracellular addition of $100 \,\mu\text{M}$ NPPB produced a strong block to 2% of its control value with an almost complete recovery after washout (79% recovery). In another neuron (Fig. 3*C*), 1 mM DIDS also blocked the Ca²⁺-activated current (2% of its control value) while the recovery was only partial (53% of control value).

On average, the current amplitude in the presence of $100 \,\mu\text{M}$ NPPB or 1 mm DIDS was, respectively, 16% or 20% of the control value (Fig. 3*B* and *C*, panels on the right). After washout of NPPB or DIDS, the current recovered on average to about 60% of its control value for both blockers (56% for NPPB and 65% for DIDS).

These results show that 100 μ M NPPB and 1 mM DIDS may also be used as efficient blockers of the Ca²⁺-activated current in olfactory sensory neurons (see also Fig. 5).

Comparison of extracellular blockers of Ca²⁺-activated currents in HEK 293T cells expressing TMEM16b/anoctamin2 and in olfactory sensory neurons

To compare the pharmacological profile of the native olfactory Ca²⁺-activated current with that of the protein that is at present the best molecular candidate for being the olfactory channel, we used the same

experimental conditions described in the previous section, uncaging caged Ca²⁺ in HEK 293T cells transfected with TMEM16b/anoctamin2.

As shown in Fig. 4A–C, each test compound partially blocked the Ca²⁺-activated current. On average, the current amplitude in the presence of 400 μ M NFA, 100 μ M NPPB or 1 mM DIDS was, respectively, 10%, 42% or 26% of the control value. After washout of the blockers, the current recovered on average to 54% (NFA), 62% (NPPB) or 64% (DIDS) of its control value.

Since the current after washout from the blockers presented some variability, to better compare the blocking efficiencies for the native and the TMEM16b/anoctamin2-induced currents, we normalized the blocked current to the average between control and washout for each experiment (Fig. 5). Ca²⁺-activated currents were reduced on average to the following percentages: 69% for NFA, 81% for NPPB, 76% for DIDS in olfactory sensory neurons, to be compared with 84% for NFA, 51% for NPPB and 70% for DIDS in HEK 293T cells transfected with TMEM16b/anoctamin2.

The blocking potencies of the three test compounds (measured both with respect to control, Figs 3 and 4, or to the average between control and washout, Fig. 5), were not significantly different in olfactory sensory neurons, while in HEK 293T cells NFA blocked more potently than NPPB.

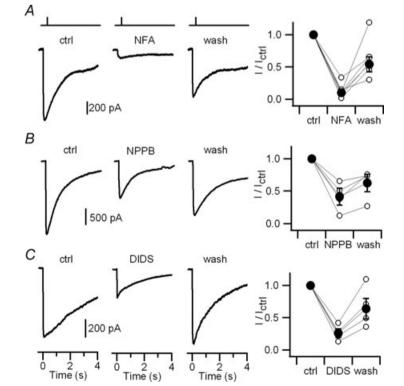


Figure 4. TMEM16b-transfected HEK 293T cells: extracellular blockage of currents activated by photorelease of caged Ca²⁺

Currents recorded in the whole-cell voltage-clamp configuration in response to photorelease of caged Ca²⁺. Ultraviolet light flashes were applied on HEK cells at the times indicated by the vertical bars in the upper panels. The holding potential was -50 mV. As in Fig. 3, the following blockers were used: A, 400 μ m NFA; B, 100 μ m NPPB; C, 1 mm DIDS. Current recordings were obtained before blocker application (control), 1–2 min after application of the indicated blockers, and 2–5 min after the removal of blockers (washout). Panels on the right show peak currents measured in the presence of each blocker and after washout, normalized to the control value before blocker application for each neuron (open circles). Average ratios are plotted as filled circles.

Anion selectivity of native Ca²⁺-activated currents in olfactory sensory neurons

To measure the ion selectivity of Ca^{2+} -activated currents in the whole-cell configuration, we photoreleased caged Ca^{2+} in the ciliary region, and recorded the current at various holding potentials while changing the ion composition in the extracellular solution. To avoid contributions from Ca^{2+} -activated K^+ currents the intracellular monovalent cation was Cs^+ .

In a first set of experiments, the intracellular and extracellular Cl⁻ concentrations were very similar ([Cl⁻]_o = 149 mM, [Cl⁻]_i = 143 mM) and the calculated equilibrium potential for Cl⁻ was -1.0 mV. Figure 6A shows currents in response to Ca²⁺ concentration jumps obtained by photorelease of Ca²⁺ when an olfactory sensory neuron was held at the indicated holding potentials from -15 to +15 mV. The rise time of the response was fast and was fitted by a single exponential with $\tau = 27$ ms at +5 mV and $\tau = 29$ ms at -5 mV.

Figure 6*B* shows current–voltage relations measured at various times after Ca²⁺ photorelease. The $V_{\rm rev}$ extrapolated from each current–voltage relation did not significantly change with time and was 0.4 mV, close to the expected Cl⁻ equilibrium potential (-1.0 mV). The average $V_{\rm rev}$ from several neurons was 1.0 ± 1.1 mV (n = 13).

Although $V_{\rm rev}$ was very close to the calculated equilibrium potential for Cl⁻, a non-selective cation current might also contribute to the Ca²⁺-activated current. If Ca²⁺-activated channels are permeable to cations, the replacement of Na⁺ by choline, a large organic cation that is usually impermeant in cation channels, should cause a large shift of the $V_{\rm rev}$ (Franciolini & Nonner,

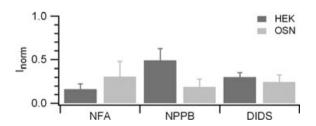


Figure 5. Comparison of extracellular blockage of Ca^{2+} -activated currents

Average ratios of currents measured in the presence of each blocker normalized to the average between control and washout currents. Experiments were from olfactory sensory neurons (OSN, n=3–8) or from HEK 293T cells transfected with TMEM16b/anoctamin2 (HEK, n=4–6). The block efficacy of different compounds was not significantly different in olfactory sensory neurons (ANOVA, F=0.33, P=0.72). In contrast in HEK 293T cells, blockage by NFA was significantly bigger from NPPB block (ANOVA, Tukey test P<0.05), but not different from DIDS block. However, current blockages in olfactory sensory neurons and in HEK 293T cells were not significantly different (unpaired t test).

1994; Qu & Hartzell, 2000; Hille, 2001). Instead, we found that the average $V_{\rm rev}$ measured in the presence of 140 mM choline chloride remained close to 0 mV, with an average value of 0.4 ± 0.6 mV (n=3; data not shown), indicating that the Ca²⁺-activated current was indeed mainly carried by Cl⁻.

To measure the selectivity among anions we replaced 140 mm NaCl in the Ringer solution with NaNO₃, NaI, NaSCN, NaBr or NaMeS and estimated the $V_{\rm rev}$ in the presence of each anion. After each anion substitution, the Ringer bathing solution still contained 9 mm Cl⁻, producing a calculated equilibrium potential for Cl⁻ of +70.2 mV.

Figure 6*C* shows recordings in the presence of nitrate Ringer solution. Surprisingly, multiple current components were clearly evident at several holding potentials. At -29 mV, a fast outward current ($\tau=15$ ms) with a peak amplitude of +45 pA, was followed by a slower current component reaching a value of 0 pA about 250 ms after the flash and -75 pA at 2.5 s, as illustrated in Fig. 6*D*.

Current–voltage relations were measured at various times after Ca^{2+} photorelease (Fig. 6E) and the $V_{\rm rev}$ extrapolated from each current–voltage relation was plotted as a function of time in Fig. 6F. The $V_{\rm rev}$ in nitrate Ringer solution showed a time dependence with a gradual shift from a value of -34 mV at 50 ms, toward a less negative value of about -24 mV during the following 2-4 s.

In the presence of NaNO₃, we measured a time-dependent shift of $V_{\rm rev}$ in each of four tested neurons. The average $V_{\rm rev}$ in the presence of NO₃⁻ was -28.6 ± 2.5 mV (n=4) for the fast current component, and -17.3 ± 3.6 mV (n=4) for the current component measured at 2–4 s. The shift of $V_{\rm rev}$ toward more negative values upon substitution of Cl⁻ with NO₃⁻ indicates that NO₃⁻ was more permeant than Cl⁻.

Substitution of chloride Ringer with iodide Ringer solution also revealed the presence of multiple current components (Fig. 7A). At -25 mV a fast outward current with a peak amplitude of +270 pA 17 ms after the flash, was followed by a slower component reaching a value of -150 pA within 2 s from flash release. At -35 mV a fast inward current component with a value of -270 pA at 50 ms was followed by an additional slower component that reached -500 pA within 2 s (Fig. 7*B*). Thus, also in iodide Ringer solution, $V_{\rm rev}$ was time dependent, with a gradual shift from -30 mV at 50 ms, to -18 mV at 2-4 s (Fig. 7*C*).

A similar time dependence of $V_{\rm rev}$ was measured in isothiocyanate Ringer solution (from -36 to -21 mV for the experiment in Fig. 7C). In the presence of bromide Ringer solution, $V_{\rm rev}$ had a small time dependence (from -17 to -14 mV for the experiment in Fig. 7C), only in 2 out of 6 olfactory sensory neurons.

The average reversal potential of the fast component was -30.5 ± 3 mV (n = 6) in the presence of I⁻, -42 ± 5.5 mV (n = 4) in SCN⁻ and -13.8 ± 2.2 mV (n = 6) for Br⁻.

When chloride Ringer was replaced by methanesulfonate Ringer solution, current recordings showed that MeS⁻ was much less permeant than Cl⁻; indeed V_{rev} was +34 mV and remained fairly constant at different times (Fig. 7*C*). The average reversal potential was +31.6 \pm 4.3 mV (n = 5). The selectivity sequence was SCN⁻ \approx I⁻ \approx NO₃⁻ > Br⁻ > Cl⁻ > MeS⁻ (Fig. 10).

These results demonstrate that the current activated by Ca^{2+} photorelease in the ciliary region of olfactory sensory neurons is anion selective and that for some anions $V_{\rm rev}$ gradually shifted with time toward less negative values.

Anion selectivity of Ca²⁺-activated currents in HEK 293T cells expressing TMEM16b/anoctamin2

To investigate whether Ca^{2+} -activated currents induced by TMEM16b/anoctamin2 in HEK 293T cells exhibits a time-dependent V_{rev} in the presence of some anions, we used the same experimental protocols as in olfactory sensory neurons.

Figure 8A shows recordings from a cell in chloride Ringer solution. As in olfactory sensory neurons, the rise time of the response to photorelease of Ca^{2+} in NaCl was fast and was well fitted by a single exponential with $\tau = 14 \, \text{ms} \, \text{at} + 20 \, \text{mV}$ and $\tau = 4 \, \text{ms} \, \text{at} - 20 \, \text{mV}$. Current–voltage relations measured at various times (Fig. 8B) showed that the estimated V_{rev} was $-0.2 \, \text{mV}$ and was not time

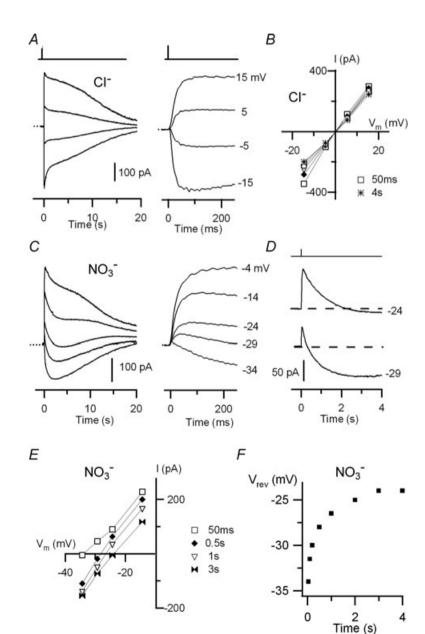


Figure 6. Olfactory sensory neurons: current responses induced by photorelease of Ca²⁺ in the presence of external Cl⁻ or NO₃⁻

Currents recorded from isolated mouse olfactory sensory neurons in the whole-cell voltage-clamp configuration in response to photorelease of caged Ca²⁺ in the cilia at the time indicated by the vertical bars at the top. A, currents from an olfactory sensory neuron were recorded in almost symmetrical CIsolutions at the indicated holding potentials and plotted on different time scales. The dotted line indicates the zero current level. B, current-voltage relations measured from recordings in A at different times from the flash: 50 ms square, 0.5 s diamond, 1 s triangle, 2 s circle, 3 s double triangle, 4 s asterisk. C, currents from another olfactory sensory neuron recorded in the presence of external NO₃⁻ at the indicated holding potentials and plotted on different time scales. The dotted line indicates the zero current level. D, recordings in external NO_3 at -24 and -29 mV are shown on a different scale to better illustrate the time dependence of V_{rev} . E_r current-voltage relations in external NO₃⁻ measured from recordings in C at different times from the flash: 50 ms open square, 0.5 s diamond, 1 s triangle, 3 s double triangle. F, V_{rev} as a function of time for external NO₃⁻ from recordings in C.

dependent. Its average value was -0.7 ± 0.7 mV (n = 21). Moreover, when NaCl was replaced by choline chloride the average $V_{\rm rev}$ was -0.5 ± 0.6 mV, n = 4 (data not shown), similar to that in NaCl, indicating that the current was mainly carried by Cl⁻, as in olfactory sensory neurons.

In the presence of nitrate Ringer solution multiple current components were evident (Fig. 8C). The current amplitude at -20 mV was +230 pA 70 ms after the flash and -18 pA at 2.5 s (Fig. 8D). Current–voltage relations as a function of time (Fig. 8E) indicate that $V_{\rm rev}$ gradually shifted with time from -29 mV to about -20 mV at 2–4 s (Fig. 8F). On average, $V_{\rm rev}$ in the presence of NO₃ $^-$ was -26.8 ± 1.5 mV (n = 4) for the fast current component, and -19.5 ± 4.8 mV (n = 4) for the current component measured at 2–4 s.

Figure 9 shows a summary of results obtained with the other anions: in iodide Ringer solution (Fig. 9A and B), at -20 mV a fast current component of +140 pA was measured at 200 ms, followed by a slower component reaching -20 pA at 5 s. A fast and a slow component were also evident at -40 mV, with a value of -100 pA at 80 ms that reached -430 pA within 4 s. The time dependence of $V_{\rm rev}$ for I⁻ is illustrated in Fig. 9C, where $V_{\rm rev}$ shifted from -26 to -18 mV at 2-4 s.

 $V_{\rm rev}$ also had a significant time-dependent shift in isothyocianate Ringer solution (from -38.5 to -18 mV for the experiment in Fig. 9C), while the shift was smaller in bromide Ringer solution (from -11.2 to -7 mV for the experiment in Fig. 9C).

The average reversal potential of the fast component was -34.9 ± 1.8 mV (n = 4) in the presence of I⁻, -29.0 ± 4.0 mV (n = 5) in SCN⁻ and -15.7 ± 0.4 mV (n = 5) in Br⁻.

Figure 9*C* shows that recordings in methanesulphonate Ringer solution exhibited a single current component with a time-independent $V_{\rm rev}$ of +27 mV. The average $V_{\rm rev}$ was +39.7 \pm 4.6 mV (n=6) confirming that MeS⁻ is less permeant than Cl⁻.

Figure 10 summarizes the average $V_{\rm rev}$ of the fast component in olfactory sensory neurons and in HEK 293T cells expressing TMEM16b/anoctamin2. For both types of currents the selectivity sequence was the following: $SCN^- \approx I^- \approx NO_3^- > Br^- > Cl^- > MeS^-$.

Furthermore, these results show that, similarly to olfactory sensory neurons, also in HEK 293T cells expressing TMEM16b/anoctamin2, the $V_{\rm rev}$ gradually shifted as a function of time toward less negative values in the presence of external ${\rm NO_3^-}$, ${\rm I^-}$, SCN⁻ and, to a lesser extent, in Br⁻, while it did not change with time in Cl⁻ and in MeS⁻.

Discussion

Expression of TMEM16b/anoctamin2 in the ciliary layer

In this study, we show by RT-PCR that TMEM16b/ anoctamin2 and some other members of the TMEM16/anoctamin family are expressed in the olfactory epithelium. Our results are in agreement with other reports showing the presence not only of TMEM16b/anoctamin2, but also of a/1 and k/10, whereas not all studies agree on other family members (Stephan et al. 2009; Rasche et al. 2010). Indeed, we identified also the presence of d/4, g/7 and j/9 that could not be detected in other reports (Rasche et al. 2010). Differences might

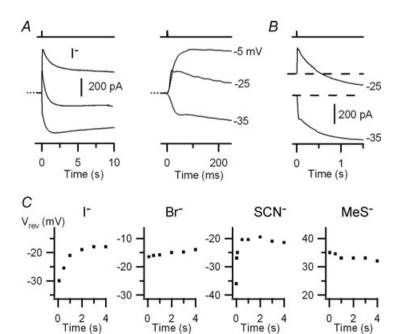


Figure 7. Olfactory sensory neurons: current responses induced by photorelease of Ca²⁺ in external I⁻, SCN⁻, Br⁻ or MeS⁻

A, currents recorded at the indicated holding potentials from an olfactory sensory neuron bathed in external iodide Ringer solution. The dotted line indicates the zero current level. B, recordings in external I^- at -25 and -35 mV are shown on a different scale to better illustrate the time dependence of $V_{\rm rev}$. C, $V_{\rm rev}$ as a function of time for external I^- (from the recordings in A), Br^- , SCN^- and MeS^- , each from a different neuron.

be due to several reasons. The discrepancy for d/4 and g/7 might reflect a higher efficiency of our amplification reaction since both are likely to be poorly expressed in the olfactory epithelium. The different result for j/9 that, in our hands, appears abundantly expressed in the olfactory epithelium, is unlikely to be caused by the recognition of a different isoform, as oligonucleotide primers employed by Rasche and colleagues are internal to ours. It remains plausible that the modality for tissue collection may reveal an important variable to explain this inconsistency, although in our study, olfactory epithelia were isolated by distinct operators without appreciable differences in the final results.

Although not all studies agree on the specific family members, at present, only the mRNA of TMEM16b/anoctamin2 has been shown to be specific to

mature olfactory sensory neurons by *in situ* hybridization (Yu *et al.* 2005), and proteomic screening of ciliary membranes (Mayer *et al.* 2009; Stephan *et al.* 2009) revealed only TMEM16b/anoctamin2 as a prominent protein in the olfactory cilia. Finally, the expression of TMEM16b/anoctamin2 in the ciliary layer of olfactory sensory neurons, has been independently confirmed by immunohistochemistry in our study, using the antibody developed by Stöhr *et al.* (2009), and in other recent reports using two other different antibodies (Hengl *et al.* 2010; Rasche *et al.* 2010).

Taken together, all these results indicate that TMEM16b/anoctamin2 is strongly expressed in the cilia of olfactory sensory neurons, where olfactory transduction takes place, while some other members of the family might be present at a lower level or in other cell types.

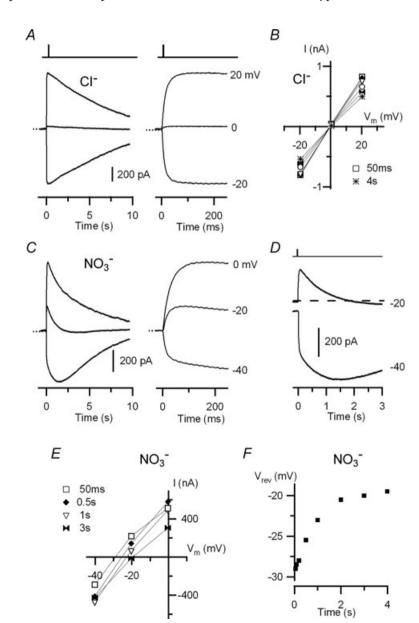


Figure 8. Anion selectivity of the TMEM16b/anoctamin2-mediated current in external CI⁻ or NO₃⁻

Currents recorded from HEK 293T cells expressing TMEM16b/anoctamin2 in the whole-cell voltage-clamp configuration in response to photorelease of caged Ca²⁺ at the time indicated by the vertical bars at the top. A, currents from a cell were recorded in almost symmetrical CI⁻ solutions at the indicated holding potentials and plotted on different time scales. The dotted line indicates the zero current level. B, current-voltage relations measured from recordings in A at different times from the flash: 50 ms square, 0.5 s diamond, 1 s triangle, 2 s circle, 3 s double triangle, 4 s asterisk. C, currents from another cell were recorded in the presence of external NO₃⁻ at the indicated holding potentials and plotted on different time scales. The dotted line indicates the zero current level. D, recordings in external NO_3 at -40 and -20 mV are shown on a different scale to better illustrate the time dependence of V_{rev} . E, current-voltage relations in external NO_3 measured from recordings in C at different times from the flash: 50 ms square, 0.5 s diamond, 1 s triangle, 3 s double triangle. F, V_{rev} as a function of time for external NO₃⁻ from the recordings in C.

Extracellular blockers

In previous studies, the pharmacological profile of the native olfactory Ca²⁺-activated Cl⁻ current has been investigated in detail only by applying Cl- channel inhibitors at the cytoplasmic side of isolated cilia from frog olfactory sensory neurons (Kleene & Gesteland, 1991; Kleene, 1993; reviewed by Frings et al. 2000). However, it is well known that some compounds have very different effects when applied to the extracellular or to the intracellular side of the membrane. The most commonly used extracellular blockers for Ca²⁺-activated Cl⁻ currents in olfactory sensory neurons are NFA (Lowe & Gold, 1993; Reisert et al. 2005; Boccaccio et al. 2006; Boccaccio & Menini, 2007; Antolin et al. 2010), which blocks also from the intracellular side (Kleene & Gesteland, 1991; Kleene, 1993; Reisert et al. 2003; Pifferi et al. 2006b), and SITS (Kurahashi & Yau, 1993; Lowe & Gold, 1993), which is a much less effective blocker from the intracellular side (Kleene & Gesteland, 1991; Pifferi et al. 2006b). We determined that the blocking properties of two other compounds, NPPB and DIDS, are side-specific with extracellular inhibition of 85% for 100 μ M NPPB and 76% for 1 mm DIDS, while Kleene and Gesteland (1991) measured a very poor intracellular inhibition: 32% for 300 μ m NPPB, and 5% for 100 μ m DIDS (see Table 2 of Frings et al. 2000).

Similar side-specific effects of NPPB and DIDS on TMEM16b-induced currents were obtained in this and in a previous study (Pifferi *et al.* 2009*a*). Moreover, we found that the average percentages of current inhibition of TMEM16b-induced currents were similar to those we measured in olfactory sensory neurons. Taken together, the

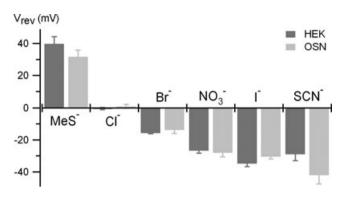


Figure 10. Comparison of anion selectivity for native and TMEM16b/anoctamin2-mediated current

Average $V_{\rm rev}$ for the fast component measured in the presence of the indicated external anions. Experiments were from olfactory sensory neurons (OSN, n=4–13) or from HEK 293T cells transfected with TMEM16b/anoctamin2 (HEK, n=4–21). $V_{\rm rev}$ in olfactory sensory neurons and in HEK 293T cells were not significantly different (unpaired t test).

results on intracellular (Kleene & Gesteland, 1991; Kleene, 1993; Pifferi *et al.* 2009*a*) and extracellular blockage of native olfactory and TMEM16b-induced currents indicate that NPPB and DIDS block the two currents in a similar side-specific manner.

It is of interest to note that in some neurons NPPB or DIDS blocked the current almost completely, while such a large inhibition was never observed in TMEM16b-induced currents in HEK cells. A small difference in results between native and expressed channels has also been pointed out by Saidu *et al.* (2010). In fact, in excised patches the native current slightly inactivates at positive

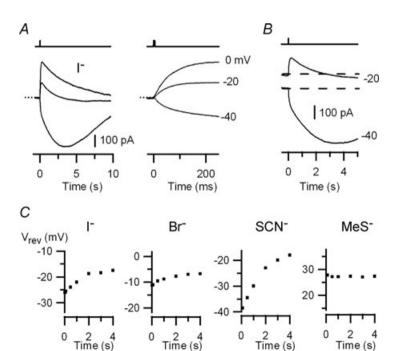


Figure 9. Anion selectivity of the TMEM16b/anoctamin2-mediated current in external I⁻, SCN⁻, Br⁻ or MeS⁻

A, currents from a HEK 293T cell expressing TMEM16b/anoctamin2 were recorded in external iodide Ringer solution at the indicated holding potentials. The dotted line indicates the zero current level. B, recordings in external I $^-$ at $^-$ 20 and $^-$ 40 mV are shown on a different scale to better illustrate the time dependence of $V_{\rm rev}$. C, $V_{\rm rev}$ as a function of time for external I $^-$ (from recordings in A), Br^- , SCN^- and MeS^- , each from a different cell.

membrane potentials (Reisert *et al.* 2003), whereas the TMEM16b/anoctamin2-induced current in HEK cells does not (Stephan *et al.* 2009; Saidu *et al.* 2010). These differences may point to the possibility that additional subunits and/or splice variants constitute the native channel, as recently suggested by Saidu *et al.* (2010).

Dynamic ion selectivity

Another important property of ion channels is ion selectivity. In previous reports, the ion selectivity of native olfactory Ca²⁺-activated Cl⁻ currents has been measured only from inside-out membrane patches excised from the knob/ciliary region of olfactory sensory neurons, exchanging ions at the intracellular side of the membrane (Reisert *et al.* 2003; Pifferi *et al.* 2006*b*). The ion selectivity of native channels in isolated olfactory sensory neurons, replacing extracellular Cl⁻ with other ions, has not been estimated yet with the exception of substitution of Cl⁻ with the largely impermeant gluconate to demonstrate that the Ca²⁺-activated current was carried by Cl⁻ (Takeuchi *et al.* 2009).

We found that some anions larger than Cl^- , such as NO_3^- , I^- , SCN^- and Br^- , were more permeant than Cl^- , while MeS^- was less permeant. Interestingly, in the presence of NO_3^- , SCN^- and I^- , a gradual time-dependent shift of V_{rev} was clearly evident. Indeed, V_{rev} in the presence of these anions changed with time, shifting about 10 mV toward less negative values over a few seconds after photorelease of Ca^{2+} , indicating that channels became less permeant to some foreign anions. To the best of our knowledge, this is the first demonstration that the native olfactory Ca^{2+} -activated Cl^- current exhibits dynamic ion selectivity.

What is the origin of the dynamic ion selectivity? A possibility is that V_{rev} changes with time because of a restricted anion diffusion: if anions accumulate at the intracellular mouth of the channel, the local concentration gradient will change modifying V_{rev} . In the narrow ciliary compartment, restricted anion diffusion might cause the accumulation of anions flowing into the cilia near the intracellular mouth of the channel, therefore leading to a different concentration gradient and a modification of V_{rev} . However, it is unlikely that this effect is the origin of the time-dependent change in V_{rev} , because such an effect is expected to be present with all anions, whereas it was not observed in the presence of Cl⁻ or of MeS⁻. Furthermore, in some experiments (Figs 6D and 7B) the current reversed direction, which should not result from restricted diffusion. In fact, if the initial NO₃⁻ or I- influx (outward current) causes an accumulation of NO₃⁻ or I⁻ inside the cilium, this should only proceed until equilibrium (zero current) and it cannot account for the Cl⁻ efflux (inward current) measured subsequently. Finally, an environmental-mediated restriction of anion diffusion, typical of the olfactory ciliary compartment, is unlikely to exist in TMEM16b/anoctamin2-transfected HEK 293T cells that, in turn, display a similar dynamic ion selectivity.

Since olfactory sensory neurons contain several types of channels, we cannot exclude that multiple current components might be due to the activation of different types of channels. Again, however, when the same experimental protocols were applied to HEK 293T cells expressing TMEM16b/anoctamin2, we obtained results very similar to those observed in the native olfactory channels. In HEK 293T cells, multiple current components were clearly evident in the presence of the same anions: NO_3^- , I^- and SCN^- , and absent with Cl^- or MeS^- . Moreover, the average $V_{\rm rev}$ for each anion was not significantly different between the two systems.

Our results have at least two important consequences: on one side they show that Ca²⁺-activated Cl⁻ currents in olfactory sensory neurons have multiple components with different anion selectivity and, on the other side they show that heterologous expression of TMEM16b/anoctamin2 in HEK 293T cells reproduced similar results, indicating that time-dependent ion selectivity is not due to different types of channels.

The presence of multiple current components is not unique to the Ca^{2+} -activated Cl^- current present in olfactory sensory neurons since, for example, it has been found in *Xenopus* oocytes (Boton *et al.* 1989; Kuruma & Hartzell, 1999). More recently, Schroeder *et al.* (2008) clearly showed the presence of multiple current components with different anion selectivity, activating Ca^{2+} -activated Cl^- currents in *Xenopus* oocytes by photorelease of caged IP_3 . Moreover, Schroeder *et al.* (2008) demonstrated that the TMEM16a/anoctamin1 channel expressed in *Axolotl* oocytes, which do not have endogenous Ca^{2+} -activated Cl^- currents, also exhibited current components with different V_{rev} , suggesting that the multiple components originated from different states of the same channel.

Furthermore, it is of interest to note that dynamic changes in ion selectivity are not a peculiarity of anion channels, and they have been revealed in cation channels (Khakh & Lester, 1999). For example, the P2X receptor of mast cells allows a time-dependent membrane permeability to large molecules (Cockcroft & Gomperts, 1979), the transient receptor potential vanilloid 1 (TRPV1) channel shows a time- and concentration-dependent change in ion selectivity in the presence of prolonged exposure to chemical agonists (Chung *et al.* 2008), and a mutant *N*-methyl-D-aspartate (NMDA) channel exhibits multiple current components with time-dependent ion selectivity (Schneggenburger & Ascher, 1997). Single-channel analysis of the mutant NMDA channel revealed the existence of at least two

subconductance states with different ion selectivity, indicating a strong coupling between permeation and gating (Schneggenburger & Ascher, 1997). Zheng and Sigworth (1997) also showed by single-channel analysis that a mutant *Shaker* K⁺ channel exhibits two subconductance states with different ion selectivity.

Another example of coupling between gating and permeation occurs in the cyclic nucleotide-gated channel in retinal rods. Indeed, it has been shown that in intact rod photoreceptors, selectivity among divalent cations changes with different levels of cGMP (Cervetto *et al.* 1988), and that in inside-out patches from photoreceptors, both the channel open probability and the selectivity of Ca²⁺ over Na⁺ increases with cGMP concentration (Hackos & Korenbrot, 1999). In addition, the linkage between selectivity and gating is specific for divalent cations, whereas it does not occur if only monovalent cations are present (Hackos & Korenbrot, 1999).

Previous studies on Ca²⁺-activated Cl⁻ currents from various cell types showed that some foreign anions at the extracellular side affect gating by modifying channel kinetics (Evans & Marty, 1986; Greenwood & Large, 1999; Perez-Cornejo *et al.* 2004). Greenwood and Large (1999) suggested that some permeant anions might modulate the kinetics of Ca²⁺-activated Cl⁻ channels in smooth muscle cells by binding to a site located on the external surface of the channel, which may be part of the channel representing the selectivity filter.

Our experiments were obtained by activating channels by a fast jump in Ca^{2+} concentration that decreased with time by diffusion and/or by active extrusion. In the presence of some anions, we observed a time-dependent shift of V_{rev} toward less negative values, corresponding to a decreased ion selectivity. A possible mechanism explaining the dynamic ion selectivity of native olfactory and TMEM16b/anoctamin2-induced Ca^{2+} -activated Cl^- currents is the presence of at least two open states with different ion selectivity and Ca^{2+} -dependent open probability. Indeed, our results are consistent with a model in which the more selective open state is favoured by high Ca^{2+} concentrations and the less selective open state by low Ca^{2+} concentrations.

The understanding of this phenomenon at the molecular level by future mutational and structural analyses, will clarify the molecular mechanisms of gating and permeation of Ca²⁺-activated Cl⁻ channels, contributing to increase the knowledge about their functioning in physiological and pathophysiological processes.

Conclusions

In conclusion, we confirmed by immunohistochemistry that TMEM16b/anoctamin2 is expressed in the

ciliary layer, and showed that TMEM16b/anoctamin2 reproduced the phenotypes of the native olfactory Ca²⁺-activated Cl⁻ currents, including the timedependent change in selectivity. Taken together with previous studies on inside-out patches (Pifferi et al. 2009a; Stephan et al. 2009), these results contribute to strongly indicate that TMEM16b/anoctamin2 is likely to be the major subunit of the native olfactory Ca²⁺-activated Cl⁻ current. Future studies should examine the presence of olfactory Ca²⁺-activated Cl⁻ currents in mice in which the TMEM16b/anoctamin2 gene is deleted. Moreover, the combination of molecular biology studies and functional measurements will clarify if additional subunits and/or splice variants belonging to the TMEM16/anoctamin or to other protein families are also part of the native Ca²⁺-activated Cl⁻ channel.

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Author contributions

All authors contributed to the conception and design of the experiments and approved the final version of the manuscript. RT-PCR experiments were performed by G.M. and R.T. at the University of Parma. Immunohistochemistry was done by M.D. and electrophysiological experiments were performed by C.S. and A.B. at SISSA, Trieste.

Acknowledgements

We thank H. Zhao (The Johns Hopkins University School of Medicine, Baltimore, MD, USA) for kindly providing the mouse olfactory TMEM16b/anoctamin2 DNA construct; H. Stöhr (Universitat Regensburg, Regensburg, Germany) for the gift of the TMEM16b monoclonal antibody; S. Pifferi, F. Celsi and all members of the laboratory for discussions. This study was supported by grants from the Italian Ministry of Education, University and Research (MIUR) and from the Italian Institute of Technology.

4 DISCUSSION

The Ca²⁺-activated Cl⁻ current in the cilia of olfactory sensory neurons accounts for most of the depolarizing current in olfactory transduction (Boccaccio & Menini, 2007; Kleene, 2008). Despite its important role, the molecular identity of olfactory Ca²⁺-activated Cl⁻ current is not definitely established. Bestrophin2 (Pifferi *et al.*, 2006*b*; Klimmeck *et al.*, 2009) and TMEM16b (Yu *et al.*, 2005; Mayer *et al.*, 2009; Stephan *et al.*, 2009; Hengl *et al.*, 2010; Rasche *et al.*, 2010) are expressed in the cilia of mature olfactory sensory neurons and they have been shown to induce Ca²⁺-activated Cl⁻ conductance when expressed in heterologous systems (Qu & Hartzell, 2004; Qu *et al.*, 2004; Pifferi *et al.*, 2006*b*; Schroeder *et al.*, 2008; Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009; Stöhr *et al.*, 2009). Both bestrophin2 and TMEM16b have been proposed to be a part of the Ca²⁺-activated Cl⁻ channel involved in olfactory transduction.

In the first part of this Thesis we presented a comparison between the Ca²⁺-activated Cl⁻ current in isolated olfactory sensory neurons from wild type and bestrophin2 knockout mice (Pifferi *et al.*, 2009*b*). In the second part of this Thesis we presented a comparison between the electrophysiological properties of the native Cl⁻ conductance and the Ca²⁺-activated Cl⁻ currents induced by heterologous expression of TMEM16b olfactory splice variant (Sagheddu *et al.*, 2010).

Comparison between OSNs Ca²⁺-activated Cl⁻ currents from wild type and bestrophin2 knockout mice.

Bestrophin2 protein is specifically localized in the cilia of mature olfactory sensory neurons (Pifferi *et al.*, 2006*b*; Klimmeck *et al.*, 2009). It has been indicated by Pifferi *et al.* (2006*b*) as a candidate for the Cl⁻ channel involved in olfactory transduction; indeed the Ca²⁺-activated Cl⁻ currents of native olfactory channels and of bestrophin2 expressed in HEK 293 cells showed the same anion permeability sequence, current-voltage relations quite close to linearity, voltage-independent and side-specific blockage by NFA and SITS. Nonetheless for bestrophin2 currents $K_{1/2}$ was 0.4 μ M, whereas for native currents $K_{1/2}$ was 4.7 μ M showing a difference of one order of magnitude from the native current (Pifferi *et al.*, 2006*b*).

We analyzed more in detail the role of bestrophin 2 in olfactory transduction comparing wild type and knockout mice for this protein (Pifferi *et al.*, 2009*a*). We obtained similar EOG recordings in olfactory epithelium from wild type and knockout mice, indicating that the lack of bestrophin2 does not produce a big impairment in the odorant sensitivity. Ca²⁺-activated Cl⁻ currents measured by using photolysis of caged 8-Br-cAMP or of caged Ca²⁺ within the cilia of isolated olfactory sensory neurons from wild type and knockout mice did not show significant differences. Finally, a Ca²⁺-activated Cl⁻ current was still present in excised inside-out patches from knob/cilia of olfactory sensory neurons of knockout mice. From these results we concluded that the absence of bestrophin2 in the ciliary layer of the olfactory epithelium does not significantly alter the olfactory transduction, although the possibility that some compensatory mechanisms occur in bestrophin2 knockout mice cannot be excluded.

Bestrophin2 expression pattern in olfactory sensory neurons have been recently investigated by Klimmeck *et al.* (2009). In adult mice, bestrophin2 expression has been specifically found in the cilia of mature olfactory sensory neurons, whereas in postnatal day 1 (P1) mice, bestrophin2 expression has been detected at all subcellular levels in developing neurons. Therefore Klimmeck *et al.* (2009) suggested that bestrophin2 has a role in neurogenesis, maybe as a volume-regulated anion channel contributing to the coordinated extension of cell volume in developing sensory neurons.

Bestrophin2 may also have a role in the maintenance of the intraciliary ionic homeostasis since bestrophin2 and other bestrophins have been shown to be activated by osmotic cell swelling even in the absence of Ca²⁺, indicating that they may be cell volume regulators (Fischmeister & Hartzell, 2005; Chien & Hartzell, 2007). Therefore the presence of bestrophin2 protein in the cilia of mature olfactory sensory neurons has been definitely confirmed (Pifferi *et al.*, 2006*b*; Klimmeck *et al.*, 2009; Pifferi *et al.*, 2009*b*), but further studies are required to better define its role in physiology of olfactory epithelium.

Comparison of functional properties of Ca²⁺-activated Cl⁻ currents from OSNs and TMEM16b expressed in HEK 293 cells.

TMEM16b, a member of the recently characterized protein family TMEM16/Anoctamin, produces Ca²⁺-activated Cl⁻ currents when expressed in different cell types (Schroeder *et al.*, 2008; Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009). TMEM16b mRNA has been shown to be specifically expressed in mature olfactory sensory neurons by in situ hybridization (Yu *et al.*, 2005; Hengl *et al.*, 2010); moreover proteomic screenings of ciliary membranes (Stephan *et al.*, 2009; Mayer *et al.*, 2009; Rasche *et al.*, 2010) and immunohistochemistry experiments (Rasche *et al.*, 2010; Hengl *et al.*, 2010) revealed that TMEM16b protein is localized in the olfactory cilia.

It has been shown that some electrophysiological properties of Ca²⁺-activated Cl⁻ currents induced by mTMEM16b expressed in HEK 293 and of native currents in the cilia of olfactory sensory neurons are remarkably similar. Currents in olfactory cilia (Kleene & Gesteland, 1991; Kleene, 1993*b*; Kurahashi & Yau, 1993, Reisert *et al.*, 2003; Pifferi *et al.*, 2006*b*; reviewed by Kleene, 2008) and induced by TMEM16b (Pifferi *et al.*, 2009*a*; Stephan *et al.*, 2009) showed similar sensitivity to Ca²⁺ and other divalent cations, similar small channel conductance (~1 pS), similar current-voltage relationship at saturating concentration of Ca²⁺ and the same permeability sequence.

At present TMEM16b protein is the best molecular candidate for the Ca²⁺-activated Cl⁻current in olfactory transduction.

In this Thesis we provided a further comparison between the electrophysiological properties of the Ca²⁺-activated Cl⁻ current in isolated olfactory sensory neurons and in

HEK 293 transfected with mTMEM16b. We used the whole-cell voltage clamp technique to record currents elicited by flash photolysis of caged calcium in the cilia of olfactory sensory neurons and in HEK 293 cells heterologously expressing the olfactory splice variant (Stephan *et al.*, 2009) of mTMEM16b.

The presence of TMEM16b at the surface of the olfactory epithelium was confirmed in immunohistochemistry experiments by using an anti-TMEM16b antibody (Stöhr *et al.*, 2009). In our pharmacology experiments the native current and the current induced by TMEM16b are strongly inhibited by extracellular application of NFA, DIDS and NPPB. In OSNs we confirmed extracellular blockage by NFA and showed extracellular blocking effects for NPPB and DIDS, two compounds that have been shown to be poorly effective when applied to the intracellular side of the membrane (Kleene & Gesteland, 1991). Similar side-specific effects of NPPB and DIDS on TMEM16b currents were obtained also in a previous study (Pifferi *et al.*, 2009*a*).

Ion selectivity of olfactory and mTMEM16b channels was determined by exchanging anions at the extracellular side of the membrane. We obtained for both channels the same permeability sequence, for some anions we also observed a gradual time-dependent shift of V_{rev} toward less negative values, indicating a decreased permeation through the channel. The finding that both native channel and TMEM16b display similar dynamic ion selectivity is a further point supporting the hypothesis that TMEM16b is a major subunit of the ciliary Ca^{2+} -activated Cl^- channel.

Conclusions

The molecular identity of the Ca²⁺-activated Cl⁻ channel involved in olfactory transduction has not been definitely established. Bestrophin2 and TMEM16b proteins are expressed in the cilia of OSNs and they have been proposed as candidates for the olfactory Ca²⁺-activated Cl⁻ channel. In this Thesis we showed that olfactory OSNs from bestrophin2 knockout mice still have Ca²⁺-dependent Cl⁻ currents in the olfactory cilia and they do not show significant impairments in the electrical properties of olfactory transduction; therefore the role of bestrophin2 in the cilia of OSNs remains to be further investigated. We found specific similarities between the electrophysiological properties

of the native olfactory channel and TMEM16b, supporting the hypothesis that TMEM16b is likely to be the main molecular component of the Ca²⁺-activated Cl⁻ channel.

Future studies should examine the presence of olfactory Ca²⁺-activated Cl⁻ currents in mice in which the TMEM16b/anoctamin2 gene is deleted. Moreover, the combination of molecular biology studies and functional measurements will clarify if additional subunits and/or splice variants belonging to the TMEM16/Anoctamin or to other protein families are also part of the native Ca²⁺-activated Cl⁻ channel.

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Acknowledgments

I would like to first express my gratitude to my woman guides.

To my supervisor Anna Menini for her faith in this project, for her constant, patient and expert guidance. She highlighted for me the importance of devoted attitude in our extraordinary job and I have learnt very much from working with her.

To Anna Boccaccio, who have taught me a lot with patience, for her support and perceptive and enlightening advices.

I feel very much indebted to Michele Dibattista for his daily help and discussions to improve my knowledge in the field; to Valentina Cenedese for her comments, and more generally for exploring with me the boundaries of professional friendship; to Fulvio Celsi for the discussions which helped me develop the ideas, in challenging me with alternative views; to several other colleagues, technicians and friends at SISSA, who have assisted me one way or another.

I am particularly grateful to my parents, Andrea, Sergio, Tore and especially Roberta, even though they are far away they have always trust me.

Also, my in-laws and my long standing friends support meant a great deal to me.

I am very thankful to my friends Marilena and Assunta, who have shared their ideas, free time and emotions with me over the years.

And most importantly I owe much to Giuseppe, his love in full support of my research has been vital energy for me and it is the best hope for our future together.

It would not have been possible without your help.

Thank you all,

Claudia