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CA²⁺ MICRODOMAINS IN OLFACTORY CILIA SUPPORT LOW SIGNALLING AMPLIFICATION OF ODOR TRANSDUCTION

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Abstract

Signal transduction depends critically on the spatial localization of protein constituents. A key question in odor transduction is whether chemotransduction proteins organize into discrete molecular complexes throughout olfactory cilia or distribute homogeneously along the ciliary membrane. Our recordings of Ca²⁺ changes in individual cilia with unprecedented spatial and temporal resolution using two-photon microscopy provide solid evidence for Ca²⁺ microdomains (transducisomes). Dissociated frog olfactory neurons were preloaded with caged-cAMP and Fluo-4 Ca²⁺ indicator. Ca²⁺ influx through cyclic nucleotide-gated (CNG) channels was evoked by uniformly photoreleasing cAMP, while measuring ciliary Ca²⁺. Discrete fluorescence events were clearly resolved. Events were missing in the absence of external Ca²⁺, consistent with the absence of internal Ca²⁺ sources. Fluorescence events at individual microdomains resembled single-CNG channel fluctuations in shape, mean duration and kinetics, indicating that transducisomes typically contain 1–3 CNG channels. Inhibiting the Na⁺/Ca²⁺ exchanger or the Ca²⁺-ATPase prolonged the decay of evoked intraciliary Ca²⁺ transients, supporting the participation of both transporters in ciliary Ca²⁺ clearance and suggesting that both molecules localize close to the CNG. Chemosensory transducisomes provide a physical basis for the low amplification and for the linearity of odor responses at low odor concentrations.

Keywords

Calcium; microdomains; Ca²⁺ fluorescence; odor transduction; olfactory neuron; cyclic nucleotide-gated channels

INTRODUCTION

Odor transduction takes place in the cilia of the sensory neurons of the olfactory epithelium. Odor binding to specific receptors triggers a G_{olf}-mediated cascade where activation of adenylyl cyclase III (ACIII) raises cAMP concentration. This nucleotide directly gates ciliary cyclic nucleotide-gated (CNG) channels (Schild and Restrepo, 1998), mediating the influx of Ca²⁺ ions that elicit the opening of Ca²⁺-activated Cl⁻ (Cl_{Ca}) channels. Since Cl⁻ concentration is higher in the ciliary lumen compared to the external mucus, both channels contribute to cell depolarization (Kurahashi and Yau, 1993; Lowe and Gold, 1993).

Intracellular Ca^{2+} also plays a key role in olfactory adaptation, largely by decreasing CNG channel affinity for cAMP (Chen and Yau, 1994).

Olfactory cilia ($<0.2 \mu\text{m}$ diameter, 5 to a few hundred μm long, depending on the species) are devoid of internal organelles (Menco, 1992), making the extracellular milieu the only source and sink of Ca^{2+} . In the ciliary membrane, a $\text{Na}^+/\text{Ca}^{2+}$ exchanger (NCX) (Reisert and Matthews, 1998) and a Ca^{2+} -ATPase (PMCA) (Castillo et al., 2007; Antolin et al., 2010) are the transporters responsible for Ca^{2+} clearance. It is possible that Ca^{2+} increases are confined to the local microenvironment around individual CNG channels, as with other Ca^{2+} -permeant channels (Naraghi and Neher, 1997; Fakler and Adelman, 2008). Takeuchi and Kurahashi (2008) measured electrical responses of similar amplitude to identical cAMP increments evoked in separate micrometer long segments of individual olfactory cilia, arriving to the conclusion that there was no interaction between adjacent segments of that size. On the other hand, Bhandawat et al. (2005) reported evidence that odor responses are built of small elementary units, consistent with the notion that odor transduction is organized into small protein ensembles involving a remarkably low gain. Here we directly test for the presence of the hypothetical submicrometer ciliary Ca^{2+} microdomains, whose spatial organization and relative location might be relevant for the dynamics and amplification of signal transduction.

MATERIALS AND METHODS

Isolation of olfactory sensory neurons

Adult frogs (*Rana pipiens*) were anesthetized by cooling on ice, killed by decapitation and pithed. All procedures were approved by the animal care and use committee of the University of Colorado Denver. Olfactory epithelia were removed, cut into small pieces ($\sim 1 \text{mm}^2$) and maintained for less than 8 h at 4°C in frog Ringer's solution. To dissociate the cells, pieces of tissue were torn apart with fine tweezers, passed 3 times through a fire-polished Pasteur pipette and filtered through a $70 \mu\text{m}$ mesh (Falcon). The cells were allowed to settle on a number 0 coverglass coated with CellTAK (BD Biosciences) to attach the cells, in order to avoid movement of the cilia.

Solutions

Normal Ringer's (in mM): 115 NaCl, 2.5 KCl, 2 CaCl_2 , 0.4 MgCl_2 , 10 HEPES, 10 glucose, pH 7.5; 0- Ca^{2+} Ringer's: 115 NaCl, 2.5 KCl, 0.4 MgCl_2 , 10 HEPES, 1 EGTA (no Ca^{2+} added), 10 glucose, pH 7.5; Li^+ -Ringer's, 115 LiCl, 2.5 KCl, 2 CaCl_2 , 0.4 MgCl_2 , 10 HEPES, 10 glucose, pH 7.5. The PMCA blocker 5-(and-6)-carboxyeosin diacetate, succinimidyl ester (CE, 50 μM ; Molecular Probes, Eugene OR) or the PDE inhibitor 3-Isobutyl-1-methylxanthine (IBMX, 1 mM) were added to Normal Ringer's, when needed. Chemicals were purchased from Sigma Chemical (St. Louis, MO), unless otherwise specified.

Calcium measurements in olfactory cilia

Cells were preincubated at room temperature for 40 min in Ringer's solution supplemented with 10 μM of Fluo-4 acetoxymethyl ester probe (Fluo-4 AM, Molecular Probes) and then washed twice (10 min each) with Ringer's solution. Cells were then incubated for at least 25 min previous to the experiment in extracellular solution supplemented with 250 μM of the cell-permeable caged-cAMP (4,5-dimethoxy-2-nitrobenzyl adenosine 3',5'-cyclicmonophosphate, Molecular Probes). Cilia presenting a straight segment of at least 5 μm were chosen for Ca^{2+} fluorescence measurements under line raster scan mode.

All measurements were made with a two photon microscope (Zeiss, model LSM-510 NLO), equipped with a Ti:sapphire laser running at 795 nm, and were visualized with a 40X or 63X water immersion objective. Fluo-4 emission was collected using 500-550 nm bandpass filters.

To uncage cAMP, we placed an optic fiber (250 μm diameter) transmitting UV light flashes (0.5 to 1 ms) delivered from a Xenon flash lamp system (JML-C2, Rapp Optoelectronic) directly under the coverglass that held the cells. The beam from the optic fiber was centered within the focal field, where the cell was positioned. A scan line was positioned along a straight segment of a cilium and scanned every 3 ms or less during 1 to 3 min, providing measurement of Ca^{2+} fluorescence changes as function of time and distance in a ciliary segment. Time resolution was similar, but not identical throughout experiments, because it depended on the length of the segment examined, being higher for shorter segments than for longer ones (range 1.9 – 3.0 ms). The time resolution of our fluorescent recordings is comparable to that of previous studies on Ca^{2+} -permeant channels using the same fluorescent probe, under TIRF microscopy (Shuai and Parker, 2005; Demuro and Parker, 2006). To test the possibility that transient fluorescence events could be artifact of non-biological origin, we conducted line-scan fluorescence measurements in external solution containing the fluorophore (in its cell-impermeant form) at a concentration (10 μM) that would result in emission of light with intensities in the same range as the fluorescence measured from the cells) and three different Ca^{2+} concentrations. These measurements gave a rather uniform fluorescence signal, completely devoid of transient events, strongly supporting the biological origin of the transient fluorescence signals recorded from the cilia.

Analysis of Ca^{2+} fluorescence measurements

The time course of transient Ca^{2+} signals was quantified over a particular region of the recording where a Ca^{2+} signal was evident. Normalized fluorescence ($\Delta F/F_i$) vs. time was plotted (Fig 1D, H and Fig. 2D and H). F_i was the basal average fluorescence intensity in the same ciliary segment before exposure of the cilium to the UV light flash, while ΔF was the fluorescence intensity at any time point minus F_i . The time at which the Ca^{2+} signal declined to half its peak value ($\tau_{0.5}$) was measured for each experimental situation.

To characterize the individual fluorescent events, a small zone where Ca^{2+} increments occurred was chosen (5 pixels, $\sim 0.15 \mu\text{m}$ of cilium) and the events were quantified before and after cAMP release. In order to differentiate significant Ca^{2+} signals from basal noise, a discriminating line was defined as 2 standard deviations above the average value of the basal fluorescence preceding CNG channels activation. There was no ambiguity on sorting out the individual fluorescence events, because the magnitude of the Ca^{2+} signals was generally much higher than the baseline noise. Event durations and their spatial dimensions were measured and their cumulative distribution histograms plotted (Fig. 3). Additionally, histograms of fluorescence amplitudes of the events occurring along the cilium segment at a given time as well as those taking place at a given spot on the cilium for the duration of the experiment were built for all experimental conditions. Average duration of individual fluorescence events were expressed as mean \pm S.D. Differences between average values were obtained with Sigmaplot 10 (Systat Software Inc, Point Richmond, CA, USA).

RESULTS

We addressed the question whether transduction-dependent Ca^{2+} increases in olfactory cilia take place in microdomains by examining whether discrete Ca^{2+} fluorescence increments are elicited by sudden uniform cAMP increments in olfactory cilia. We used two-photon microscopy, which allows measuring the spatio-temporal distribution of Ca^{2+} fluorescence increments with sub-micrometer and sub-millisecond resolution, with limited contribution

from out of focus fluorescence and negligible photobleaching, contrary to conventional confocal microscopy (Denk et al., 1995; Denk and Svoboda, 1997). Olfactory neurons were preloaded with caged-cAMP and the Ca^{2+} indicator Fluo-4. cAMP was photoreleased by a UV flash. Figure 1 illustrates representative experiments conducted under 2 mM (control; A-C) and in no external Ca^{2+} (D-F). Line-scan fluorescence measurements were made every 3 ms or less for 180 s, in straight segments of individual cilia (Figs. 1A, D, red arrows). Multiple fluorescence spots were observed (Fig. 1B) along the cilium segment (ordinate, d) during the recording time under control solution (abscissa, t), indicating prominent Ca^{2+} increments at well defined spots along the cilium (see below). This is shown in further detail for different times along the experiment in the insets underneath the continuous fluorescence recording (Fig. 1B). In the experiment in Figure 1, the relative fluorescence measured in a $\sim 0.4 \mu\text{m}$ long stretch of the cilium (Fig. 1B, black arrowhead) over the whole time of the experiment incremented following the UV flash, reached a peak at ~ 35 s, subsequently decaying to basal level, with a half maximal time of 11 s (Fig 1C; $\tau_{0.5} = 10.5 \pm 1.9$ s, mean \pm S.E.M.; $n = 11$ of 17 cells tested). A similar experiment performed with cilia bathed in 0- Ca^{2+} Ringer's solution showed only a few fluorescence events (Fig. 1E,F; $n = 8$), as expected because external Ca^{2+} is the origin of intraciliary Ca^{2+} increments. Thus, we refer to the cAMP-induced discrete fluorescence increases as ' Ca^{2+} microdomains'. We noticed the existence of a basal frequency of fluorescence events preceding the UV flash, which may be attributed to the low spontaneous activity of the CNG channels (Kleene, 2000).

Inhibition of either of two Ca^{2+} transporters slows down the recovery of luminal Ca^{2+} level (Fig. 2), in accordance with the slackening in the relaxation of the Cl_{Ca} transduction current by these pharmacological agents (Castillo et al., 2007). In the presence of the PMCA blocker carboxyeosin (CE, 50 μM), ciliary fluorescence (Fig. 2A) increased at discrete sites with a more diffuse distribution compared to control conditions (Figs. 2B) and relaxed more slowly (Fig. 2C, $\tau_{0.5} = 34$ s; average $\tau_{0.5} = 30.18 \pm 6.4$ s, $n = 8$ of 14 cells tested). Abolishing NCX activity by substituting external Na^{+} with Li^{+} (Fig. 2D-F) resulted in more diffuse microdomains (Fig. 2E) and affected fluorescence decline similar to inhibiting PMCA (Fig. 2F; $\tau_{0.5} = 31$ s; average $\tau_{0.5} = 28.12 \pm 5.20$ s, $n = 6$ of 11 cells tested). This evidence supports the contribution of both transporters to Ca^{2+} clearance.

A close examination of the recordings revealed that the evoked Ca^{2+} response was made up of brief discrete events (Fig 1B, insets). Strikingly, inspection of such individual fluorescence events as a function of time revealed the occurrence of square fluctuations between a small number (1–3, depending on the spot) of well defined and regularly spaced amplitude levels, closely resembling single-channel events, suggesting a stereotyped organization of CNG channels in the ciliary membrane (Fig. 3A). Respective amplitude distributions are shown besides each trace. The recordings chosen for this type of analysis correspond to those in which separate events were discernible within the fluorescence activity; we obtained recordings where this was not possible due to excessive activity (not shown), which might reflect a higher spot density as a result of higher cAMP levels attained in such experiments.

Traces from spots containing one individual channel each from the three experimental conditions tested (control, PMCA and NCX inhibited) are shown in Fig. 3B. As in control conditions, levels ranged from one to three under the different treatment conditions. The average duration of the fluorescence events for the control trace was 3.7 ms. This value incremented by nearly 50% upon abolishing either PMCA or NCX ($\tau_f = 6.5$ and 6.8 ms, respectively). Average τ_f values for each condition (3.9 ± 0.5 ms, 6.7 ± 1.2 ms and 6.7 ± 1.7 ms, respectively; mean \pm S.D.; control vs. Li^{+} : $F = 8$, $t = -3.50$, $P = 0.008$; control vs. CE: $F = 8$, $t = -4.96$, $P = 0.001$; Student's test) are plotted in figure 3C. Figure 3D depicts fluorescence amplitude histograms each one built with data obtained from several cilia (see

figure legend), recorded under the different experimental conditions. Remarkably, all histograms display three equally spaced peaks, suggesting the presence of events with three well defined fluorescence amplitudes above the basal fluorescence level. We examined the spatial dimension of the fluorescence spots, for which we plotted the cumulative distribution of the events sizes (Fig. 3E). The events recorded under control condition were of a size at the limit of resolution of the microscope ($\sim 0.2 \mu\text{m}$). However, the width of the spots widened when either transporter was abolished, presenting average sizes of $0.4 \pm 0.5 \mu\text{m}$ under carboxyeosin and $0.4 \pm 0.3 \mu\text{m}$ in Li^+ solution. The cumulative distribution of the duration of the events under control condition shows an average value below the time resolution of the instrument ($< 2 \text{ ms}$), while under blockage of PMCA it incremented to $6.3 \pm 3.8 \text{ ms}$ and to $4.9 \pm 3.2 \text{ ms}$ when NCX was abolished (Fig. 3F).

DISCUSSION

In this work we determined the existence of Ca^{2+} microdomains comprising a small number of CNG channels in olfactory cilia and investigated Ca^{2+} dynamics following sudden uniform increments in intraluminal concentrations cAMP, confirming the participation of both NCX and PMCA on Ca^{2+} clearance.

The relaxation kinetics of the fluorescence increment is comparable with that previously observed with regular confocal microscopy in odor-stimulated salamander olfactory cilia ($\sim 7.7 \text{ s}$) (Leinders-Zufall et al., 1998). However, the latency of the Ca^{2+} -fluorescence events following UV-illumination in our experiments ($5\text{--}15 \text{ s}$) was much longer than that of the whole cell current induced by photoreleased cAMP ($\sim 0.1 \text{ s}$) (Takeuchi and Kurahashi, 2002; Castillo et al., 2007), its principal component being the Cl_{Ca} current. Its relaxation time ($\sim 11 \text{ s}$) was much longer than that of the current ($< 1 \text{ s}$) as well. Our experiments do not allow us to unravel the reason for the unexpectedly long latency. One possibility is that the whole cell current, mainly carried by Cl^- ions, directly depends on the free Ca^{2+} concentration *at the internal membrane surface*, which is sensed by the Cl_{Ca} channels localized immediately next to the open CNG channels in the microdomain (these can be as close as 20 nm). Thus, a large electrical signal rapidly develops as Ca^{2+} concentration reaches threshold for Cl_{Ca} channel activation, product of the high Ca^{2+} sensitivity of the Cl^- channels (nM) and the important amplification of this last transduction step. If the Ca^{2+} changes occur in the vicinity of the ion channels (within 20 nm) it does not necessarily translate into fluorescence change because of the very weak fluorescence signal it would evoke. In contrast, the latency preceding the detectable Ca^{2+} fluorescence events is much longer because in spite that cAMP rises uniformly in the entire ciliary lumen, it drops very quickly to sub-threshold levels within the miniscule microdomain compartments, which comprises most ciliary PDE molecules (Cygnar and Zhao, 2009). The strong PDE barrier would greatly limit cAMP diffusion from the luminal pool to the microdomain, where its concentration would gradually build up to eventually attain threshold for CNG activation. The time that this takes depends on the concentration of cAMP at the lumen and the potency of the PDE barrier. In the present work this latency was several seconds to tens of seconds. Interestingly, Kurahashi (1990) observed that the latency of the whole cell current evoked by introducing a large concentration of cAMP (0.5 mM) with a patch pipette positioned at base of the dendrite was much longer (1.4 s) than at the dendritic knob (0.2 s). In our experiments we are not aware of the caged-cAMP concentration attained in the cell after incubation in the caged compound ($250 \mu\text{M}$), nor the concentration of photoreleased cAMP, but they may have been much less than 0.5 mM . Accordingly, it does not seem unreasonable that the fluorescence event activity was preceded by long delays.

Regarding the relaxation of the responses to photorelease cAMP, we would expect the nucleotide to rapidly decline within the microenvironment surrounding the CNG channels

due to potent local PDE activity, explaining the relatively fast relaxation time of the Cl^- current. But the cAMP concentration in the bulk of the cilium would take a much longer time to decline, because the PDE molecules that hydrolyze it are confined to the membrane and likely became overwhelmed by abnormally large amounts of cAMP generated by the photolysis. This is manifested by the slowly developing but sustained activity of discrete fluorescent events. Eventually cAMP drops to basal levels, ending the fluorescent events.

Besides the strikingly similar appearance between the Ca^{2+} fluorescence fluctuations and the single CNG channel activity recorded electrically in patches excised from the dendritic knob, the dwell time obtained from the fluorescence measurements is close to that derived from electrical measurements ($\tau_o = \sim 1.5$ ms) (Zufall et al., 1991) with substantially higher time resolution (2.5 vs. ~ 0.3 KHz), indicating that the fluorescence events are manifestations of highly localized Ca^{2+} increments occurring as individual CNG channels open in one microdomain. The fluorescence levels indicate the number of CNG channels contained by individual microdomains. The prolongation of the dwell time occurring upon abolishment of PMCA or NCX can be explained by the retention of Ca^{2+} around the channel for a longer time, consistent with the participation of both transporters in ciliary Ca^{2+} removal.

A recent study showed that identical cAMP increments evoked in successive 1 μm segments of individual olfactory cilia induced electrical responses of similar amplitudes, leading to the conclusion that CNG channels were uniformly located along the cilium (Takeuchi and Kurahashi, 2008). However, our finding of <0.2 μm size Ca^{2+} microdomains suggest that if several of them activate within a 1 μm ciliary segment, the resulting electrical responses to identical stimuli would be alike. The existence of microdomains was questioned by Reisert and co-workers (2003), based on studies with inside-out patches excised from dendritic knob, some of which might have additionally contained cilia. From cAMP-evoked multichannel Cl^- currents recorded at various Ca^{2+} buffer concentrations, they concluded that CNG and Cl_{Ca} channels were ~ 120 nm apart, a distance incompatible with co-localization into a transducisome. However, there is no reason to expect a similar molecular organization or spatial distribution of channels in the cilia and knob membranes nor there is any evidence that the Cl_{Ca} channels in the knob are the same as those in the cilia.

The small number of CNG channels in the microdomains suggests that the odor transduction cascade involves low amplification, where the odor binding to a single receptor would lead to the opening of very few (typically 3 or less) CNG channels and a limited number of Cl_{Ca} channels surrounding each CNG in discrete sparse microdomains. This result is in agreement with the notion that at low concentration odor responses are the linear summation of multiple elementary responses of extremely small amplitude (Bhandawat et al., 2005). This observation is based on the statistical analysis of responses to brief odorant stimuli of low concentration, where the dose-response relation was evidently linear. A key consideration was that the OR- G_{olf} interaction time is brief (~ 0.1 ms), implying that an odor receptor has a low probability of activating one G_{olf} protein when it binds an odor molecule. On the other hand, as for other Ca^{2+} -conducting channels, a pronounced Ca^{2+} profile around a CNG channel limits to tens of nanometers from the mouth of this channel the distance at which Ca^{2+} concentration is sufficient for Cl_{Ca} activation (Naraghi and Neher, 1997; Bauer, 2001). Therefore, it is reasonable to postulate that the CNG and the Cl_{Ca} channels would localize to the Ca^{2+} microdomains, and that the number of Cl_{Ca} channels that can be housed and be functional within the microdomain cannot be large. Assuming that most of the area immediately surrounding a CNG where $[\text{Ca}^{2+}]$ attains threshold for Cl_{Ca} activation is occupied by Cl_{Ca} and a few transporter molecules, and a diameter of these proteins of 10 nm (Long et al., 2005; Fakler and Adelman, 2008), less than 10 Cl_{Ca} channels would fit in such area. Beyond this area, Ca^{2+} increments would be insufficient for opening Cl_{Ca} s (Fig. 4). This conclusion has an important implication regarding the characteristics of the Cl_{Ca}

channel, currently under debate (Frings et al., 2000; Kleene, 2009). As the Cl^- current associated to a single CNG channel is 3 to 4 times the cationic current (Lowe and Gold, 1993), the number of Cl_{Ca} in a microdomain must be sufficient to allow such anionic current, but it shall not exceed the quantity of channels that sterically fits in it. This number is directly related to the single Cl_{Ca} channel conductance; if its conductance were 1 pS (Reisert et al., 2003; Pifferi et al., 2006; Pifferi et al., 2009; Stephan et al., 2009) and as one CNG channel conducts 0.08 pA at -55 mV (Zufall and Firestein, 1993), 5-6 Cl_{Ca} channels would be required to conduct 0.32 pA, in agreement with the model of Fig 4. This implies that under physiological conditions an elementary response would be undetectable, and several of such responses should occur simultaneously in the cilia of an olfactory neuron to be physiologically significant. This is in agreement with the notion that vertebrate olfactory receptor neurons are incapable of responding to individual odorant molecules (Paysan and Breer, 2001). Interestingly, elements capable of sustaining macromolecular complexes exist in olfactory cilia, such as scaffolding proteins (Saavedra et al., 2008) or lipid rafts (Schreiber et al., 2000), although it is still unclear whether they play such a role.

The duration and width of the microdomains were similarly incremented when either of the two Ca^{2+} transporters present in the olfactory cilia was abolished. These observations imply that the NCX and PMCA effectively contribute to limit the diffusion of Ca^{2+} away from the CNG channels, suggesting that they localize in close proximity to the CNG channels, likely being part of the microdomain. This would assure an efficient removal of calcium ions after the transduction machinery is activated.

In conclusion, our data show that Ca^{2+} increments in olfactory cilia that result from CNG channel opening are confined to microdomains. Each microdomain comprises one to three CNG channels. It seems likely that these channels are associated with other transduction proteins into transducisomes, giving origin to elementary odor responses that add up generating the cellular odor response. In addition, the present results strongly support the involvement of both, NCX and PMCA in the extrusion of Ca^{2+} and in restricting Ca^{2+} increases in olfactory cilia.

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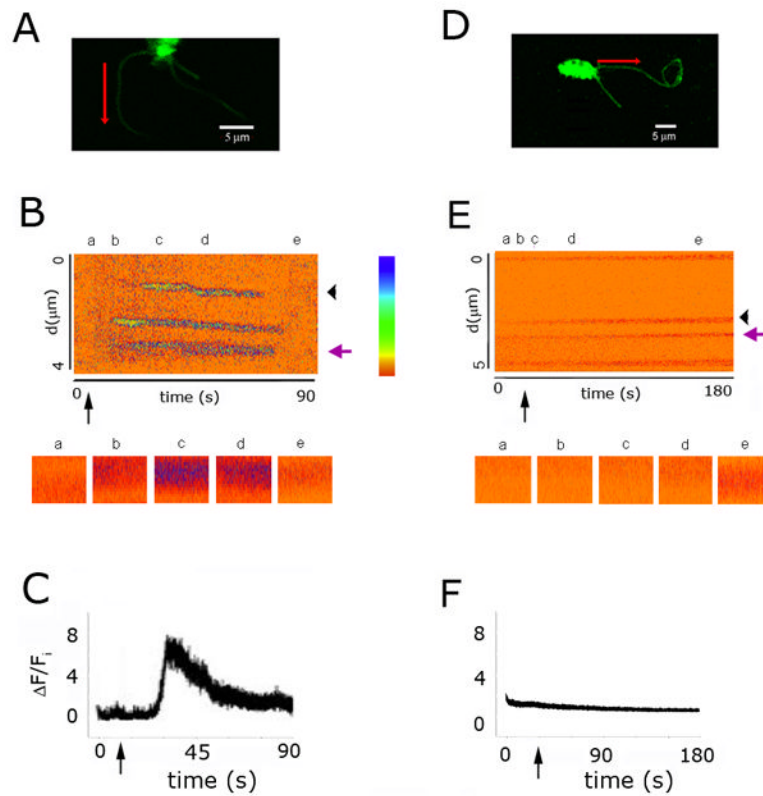


Figure 1. Localized cAMP-induced Ca^{2+} increases in olfactory cilia

A, Dendritic knob and cilia of an olfactory receptor neuron preloaded with Fluo-4 and caged-cAMP. The red arrow denotes the cilium segment from where the emission of fluorescence was collected and the direction of the scan. **B**, Ca^{2+} fluorescence changes along the cilia segment; ordinate: time, abscissa: distance; relative fluorescence intensity is color-coded, according to calibration bar. The five insets underneath the fluorescence recording show details of it at indicated the times (a-f0). Black arrow: time of the photoreleasing UV-flash. Violet arrow: ciliary segment from where insets were taken, at the times pointed out with asterisks. Black arrowhead: segment from where $\Delta F/F_i$ (**C**) was obtained. **C**, Time-course of the fluorescence change ($\Delta F/F_i$) for a selected ciliary region (black arrowhead in **B**). Relaxation half time, $\tau_{0.5} = 11$ s. **D**, Another olfactory neuron, bathed in 0-Ca^{2+} Ringer's solution. **E**, Ca^{2+} fluorescence in the ciliary segment. **F**, Relative fluorescence upon cAMP release. Labels as in **A-C**. Relative fluorescence was calculated by dividing fluorescence intensity by the average fluorescence intensity in the entire cilium for the period preceding cAMP photorelease.

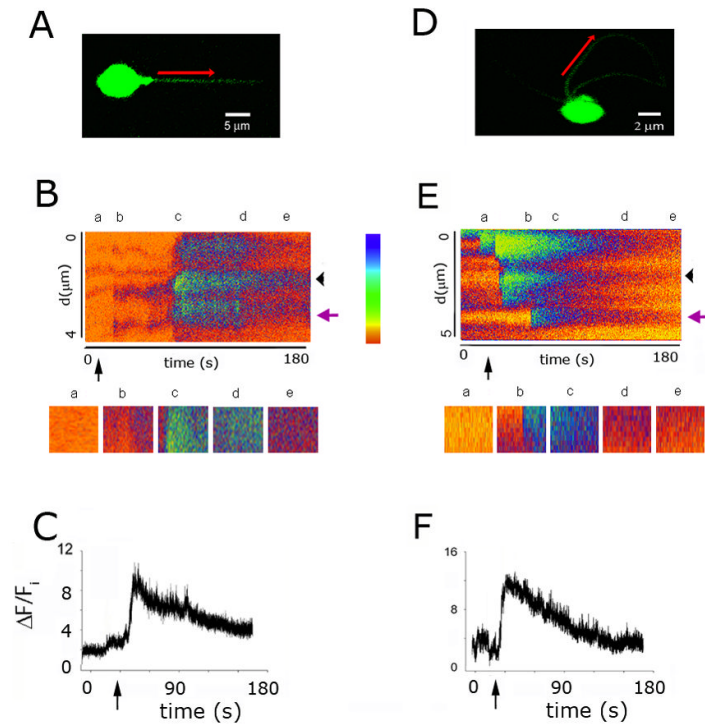


Figure 2. Effects of inhibiting PMCA and NCX on Ca^{2+} microdomains and Ca^{2+} clearance
A, Olfactory receptor neuron incubated in 50 μM carboxyeosin to inhibit PMCA, with red arrow indicating the olfactory cilia segment studied and direction of the scan. **B**, Ca^{2+} fluorescence changes along the cilia. **C**, Time-course of the fluorescence change for a selected region in the cilium. Relaxation half-time, $\tau_{0.5} = 34$ s. **D**, Ciliary segment of another olfactory neuron examined under Ringer's- Li^+ solution to abolish NCX activity. **E**, Fluorescence change as a function of distance and time. **F**, Time-course of fluorescence decay; $\tau_{0.5} = 31$ s. Labels as in Fig 1.

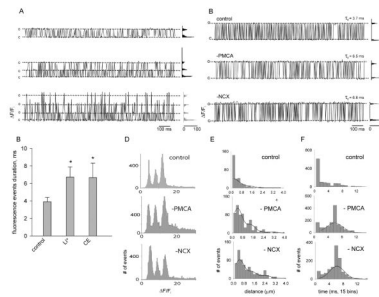


Figure 3. Unitary calcium events recorded in olfactory cilia

A, Fluorescence recordings from 3 spots, exhibiting 1, 2 and 3 amplitude levels.

Fluorescence changes were normalized by dividing the change in fluorescence (ΔF) by the average basal fluorescence before the UV flash (F_1). The 'y' scale has been arbitrarily scaled so that the change in fluorescence to the first level is the same for the three traces. **B**,

Representative high-resolution fluorescence recordings from 3 individual spots of different cilia under different condition: "Control" is normal Ringer's, "-PMCA" is Ringer's with 50 μM CE and "-NCX" is Li^+ -Ringer's. **C**, Plot showing the average durations of the fluorescence events under each condition; measurements from 5 spots each (mean \pm S.D.;

control vs. Li^+ : $F = 8$, $t = -3.50$, $P = 0.008$; control vs. CE: $F = 8$, $t = -4.96$, $P = 0.001$; Student's test) **D**, Fluorescence intensity histograms at different conditions, each one comprising spots obtained from multiple cilia. **E**, Histograms showing the cumulative

distribution of the spatial dimension of the fluorescent events induced by UV light for the different conditions, fitted with Gaussian functions. **F**. Histograms showing the cumulative

distribution of the duration of the fluorescent events under the different conditions, fitted with Gaussians functions.

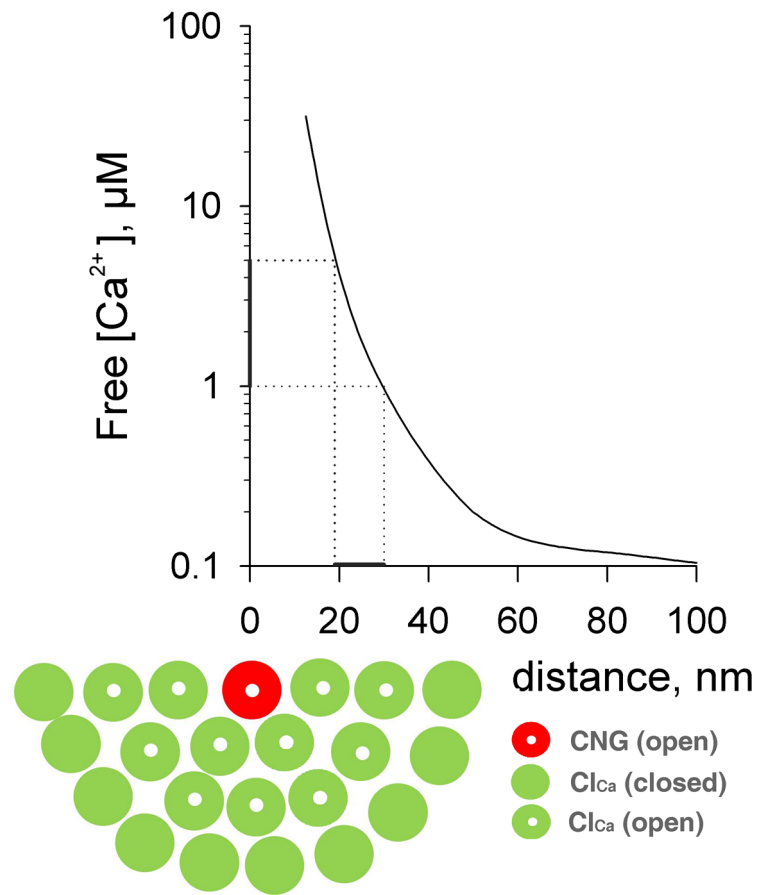


Figure 4.

Model of an olfactory cilium transducisome. The model depicts one transducisome (actually half of it, for convenience) as suggested by our results, next to the profile of free Ca^{2+} concentration as a function of distance from the mouth of a single Ca^{2+} -permeant channel, in this case a cyclic nucleotide-gated channel, adapted from (Naraghi and Neher, 1997). The CNG is surrounded by multiple Cl_{Ca} channels, whose only Ca^{2+} source is the CNG channel. The $K_{0.5}$ range of Cl_{Ca} for Ca^{2+} , 1–5 μM (Kleene and Gesteland, 1991; Reisert et al., 2003), is indicated by a bar beside the ordinate, projected to the distance axis. Proteins have been assigned an estimated size of ~10 nm (Fakler and Adelman, 2008). According to the model, $[\text{Ca}^{2+}]$ drops below threshold for Cl_{Ca} activation beyond the second row of proteins surrounding the CNG, implying that if there were further Cl_{Ca} in the transducisome, they could not open. As the Ca^{2+} transporters and protein constituents of the transduction cascade are likely to be part of the transducisome as well, the number of Cl_{Ca} within the restricted transducisome area may be even smaller.