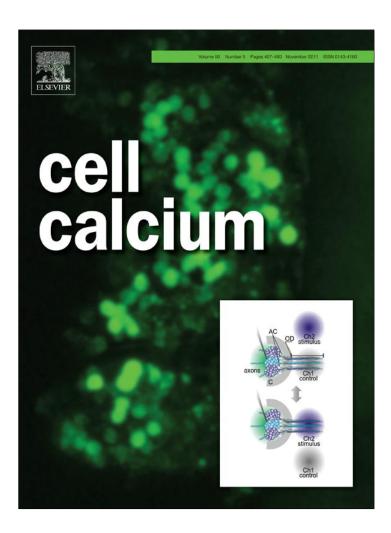
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Review

Redox regulation of calcium ion channels: Chemical and physiological aspects

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ABSTRACT

Reactive oxygen species (ROS) are increasingly recognized as second messengers in many cellular processes. While high concentrations of oxidants damage proteins, lipids and DNA, ultimately resulting in cell death, selective and reversible oxidation of key residues in proteins is a physiological mechanism that can transiently alter their activity and function. Defects in ROS producing enzymes cause disturbed immune response and disease.

Changes in the intracellular free Ca²⁺ concentration are key triggers for diverse cellular functions. Ca²⁺ homeostasis thus needs to be precisely tuned by channels, pumps, transporters and cellular buffering systems. Alterations of these key regulatory proteins by reversible or irreversible oxidation alter the physiological outcome following cell stimulation. It is therefore necessary to understand which proteins are regulated and if this regulation is relevant in a physiological- and/or pathophysiological context. Because ROS are inherently difficult to identify and to measure, we first review basic oxygen redox chemistry and methods of ROS detection with special emphasis on electron paramagnetic resonance (EPR) spectroscopy. We then focus on the present knowledge of redox regulation of Ca²⁺ permeable ion channels such as voltage-gated (CaV) Ca²⁺ channels, transient receptor potential (TRP) channels and Orai channels.

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1. Basic redox chemistry of oxygen

Many chemical processes are linked to the exchange of electrons between two or more molecular entities. The transfer of one (or more) electron(s) is associated with oxidation (loss of electron) and reduction (gain of electron) of the components. Such reactions are also known as redox reactions. The processes of reactive oxygen species (ROS) formation and elimination are exclusively of redox nature.

Molecular oxygen is the precursor of all ROS. It contains two unpaired electrons in separate (antibonding) orbitals in its outer electron sphere and is thus, by definition, a radical. Radicals have at least one unpaired electron in their orbital system and usually show high reactivity; transition metal ions also may have unpaired electrons, but are not called radicals. Although having radical character, molecular oxygen is chemically rather inert (luckily!), because high

activation energy is needed to fully reduce O_2 to water. However, single electron transfers are occurring quite readily leading to the formation of several different ROS which can be either anions or neutral molecules with quite variant reactivity. ROS with one unpaired electron are radicals, but non-radical species such as H_2O_2 are also generated and classified as ROS. The oxygen radicals generally have an unstable electronic configuration and display a high reactivity because no activation energy is required for their reaction. The most common and biologically important ROS are the superoxide radical $(O_2^{-\bullet})$ and as neutral form, the hydroperoxyl radical (HO_2^{\bullet}) , the hydroxyl radical (OH^{\bullet}) , as well as hydrogen peroxide (H_2O_2) and hypochloric acid (HOCl) (see Fig. 4).

Whether or not a redox reaction, i.e. an electron transfer, can potentially occur between two compounds (1 and 2) is depending on the effective redox potentials of both redox couples Ox1/Red1 and Ox2/Red2. The effective redox potential E of a certain redox couple is defined by the Nernst relation:

$$E = E_0 + 2.303 \frac{RT}{nF} lg \frac{[Ox]}{[Red]}$$

In the second term the factor 2.303RT/(nF) includes the temperature dependence (T, absolute temperature) and the

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number of electrons transferred (n). R and F are the gas constant (8.314 J mol $^{-1}$ K $^{-1}$) and the Faraday constant (96,487 C mol $^{-1}$), respectively (The coefficient arises from the base transformation going from the natural to the decadic logarithm). This factor is also termed Nernst factor and yields 59 mV at 25 °C or 62 mV at physiologic 37 °C for a single electron transferred (n = 1).

This second term becomes zero, when the concentrations of the oxidized and reduced form of a redox couple are equal. In that case, the effective redox potential E adopts the value of the standard reduction potential E_0 , which is measured in relation to the potential of the standard hydrogen electrode (at 25 °C, all effective concentrations at 1 mol/L and/or partial pressures of 1013 hPa for gases). The potential of the standard hydrogen electrode (SHE) is by convention defined as zero, so that the redox potential of a redox couple adopts positive or negative E_0 values depending on its electron accepting or donating properties relative to the H/H $^+$ redox couple. Because the SHE reference potential is defined in the presence of 1 mol/L of H $^+$ -ions (i.e. pH = 0), it also explicitly depends on the prevailing pH. For the biological relevant pH = 7 the reference potential at 25 °C of the SHE is given as

$$E_o^7(SHE) = E_o - 2.303 \frac{RT}{F} pH = -413 \text{ mV}$$

with $E_0 = 0 \text{ V}$ as mentioned above.

Consequently, the standard redox potentials of a redox couple involving H⁺-ions may be determined with respect to the SHE, but can be corrected for pH=7 by adding $-413 \,\mathrm{mV}$. For example, the value E_0 for the reduction of oxygen to water $(1/2O_2 + 2H^+ + 2e^- \to H_2O)$ is given as $+1.23 \,\mathrm{V}$ at standard conditions and translates to $E_0' = 0.82 \,\mathrm{V}$ corrected for pH=7. For biologic and physiologic applications usually the corrected (and primed) values E_0' of the midpoint redox potential are of relevance to predict possible redox reactions.

A redox reaction of two compounds (1 and 2) generally is written in the form

$$Red1 + Ox2 \leftrightarrow Ox1 + Red2$$
,

for which the so called half-reaction (say, of compound 1) is presented as

Red1
$$\leftrightarrow$$
 Ox1 + $ne^- + mH^+$

including explicitly the transfer of n electrons and possibly m protons.

The effective redox potential of the pair Ox1/Red1 is defined according to the Nernst equation by

$$E = E_{o} + 2.303 \frac{RT}{nF} lg \frac{[Ox1]}{[Red1]} + 2.303 m \frac{RT}{nF} lg \left[H^{+}\right]$$

Transforming to the midpoint potential E_0 ' at pH 7 and using the definition pH = $-\lg[H^+]$ we can write referring to pH 7 in biologic systems

$$E = E'_{0} + 2.303 \frac{m}{n} \frac{RT}{F} (7 - pH) + 2.303 \frac{RT}{nF} lg \frac{[Ox1]}{[Red1]}$$

Combining the constants and assuming a temperature of $37\,^{\circ}\text{C}$ we finally obtain

$$E = E'_0 + \frac{m}{n} 61.55 \,\text{mV} \,(7 - \text{pH}) + \frac{61.55 \,\text{mV}}{n} lg \frac{[\text{Ox}1]}{[\text{Red}1]} \tag{1}$$

where the effective potential E is explicitly dependent on a pH shift from a value of 7 and on the concentrations of oxidized and reduced form (with E_0 ' and E in mV). The pH-term is zero when no protons are involved, m = 0, or when the pH equals 7. It adds to the E_0 ' value, if the pH is below 7 or diminishes it for pH values above 7. For one pH unit deviation from pH 7 and for equal numbers of protons m and electrons n transferred, E_0 ' is increased or decreased

by about 62 mV. The "concentration-term" (third term in Eq. (1) above) depends on the concentrations of Ox1 and Red1 and the number of electrons *n* transferred in the redox process and is zero for equal concentrations of both components of the redox pair Ox1 and Red1.

Let us now consider an important biological half reaction, the formation of the neutral ubiquinone radical UQH• after addition of an electron and a proton to UQ

$$UQ + e^- + H^+ \leftrightarrow UQH^{\bullet}$$

for which a midpoint potential $E_0' = -40 \,\text{mV}$ is found in the literature. To visualize the pH- and concentration dependence of this redox process one can rewrite Eq. (1) with the relation $[Ox1]+[Red1]=c^*$, identifying Ox1 with UQ and Red1 with UQH• and c* being the total concentration. Then, the effective redox potential can be plotted as a function of the relative concentration of the reduced species $c_{Red1} = [Red1]/c^*$ ranging from 1 (no Ox1 present) to 0 (no Red1 present). For equal concentrations (Red1 = Ox1) the midpoint potential for different pH values can be read from the corresponding curves at $c_{Red1} = 0.5$ as indicated in Fig. 1. The E_0 '-values are shifted in positive direction for decreased pH and towards more negative values for increased pH compared to the value for pH 7. When the portion of Red1 drops to 10% $(c_{Red1} = 0.1)$ of the total concentration, the effective potential E for pH 7.4, for example, increases from $-65 \,\mathrm{mV}$ to $-5.9 \,\mathrm{mV}$ as can be inferred from the intersection of the vertical line at $c_{Red1} = 0.1$ with the curve (Fig. 1A). If the fraction of Red1 amounts to only 0.1% $(c_{Red} = 10^{-3})$ then E increases even further to +120 mV. This short example demonstrates that significant shifts in redox potential have to be considered for low concentrations of the reduced form Red1. It is also obvious from the graphs that E is decreasing massively as soon as the portion of Red1 increases and Ox1 decreases proportionally (vertical line at $c_{Red1} = 0.9$ in Fig. 1A). Such situations where one part of a redox couple is present only in minor concentrations are often found in biologicaly relevant redox reactions.

As an example, let us analyse a common biological reaction, the formation of the semiquinone radical from superoxide, in more detail, which can be described again by the formal redox equation

$$Red1 + Ox2 \leftrightarrow Ox1 + Red2$$

and explicitly by

$$O_2^- + UQ \leftrightarrow O_2 + UQH^\bullet$$

This reaction can spontaneously proceed from the left to the right side, if the redox potential of the oxidant UQ/UQH• is more positive than that of the reductant O_2/O_2 • (i.e. the free energy $\Delta G < 0$). The midpoint potential E_0 of both redox pairs (for equimolar concentrations of Ox/Red) are given in tables as $E_0'(UQ/UQH^{\bullet}) = -65 \text{ mV}$ and $E_0'(O_2/O_2^{\bullet}) = -170 \text{ mV}$, so that electrons can principally be transferred from superoxide to UQ to form semiquinone radicals until an equilibrium is reached. This can be visualized in Fig. 1B by an arrow parallel to the x-axis starting from the midpoint potential of $O_2/O_2^{-\bullet}$ which intersects with the curve of UQ/UQH $^{\bullet}$ at about $c_{Red} = 0.98$. Hence, when we have equimolar amounts of $O_2/O_2^{-\bullet}$ (and E = -170 mV) then 98% of UQ will be reduced to the semiquinone radical (right arrow in Fig. 1B). Of course we can reverse the arrow, which then means that we need 98% of reduced UQ to drive the reaction such that half of the oxygen is converted to superoxide, and we need an even higher portion of UQH* to increase the concentration of the superoxide radical

For a more realistic situation with a much smaller relative concentration of superoxide (3×10^{-3}) we will have about 13% of UQH• (left arrow in Fig. 1B), or vice versa, we need at least that amount

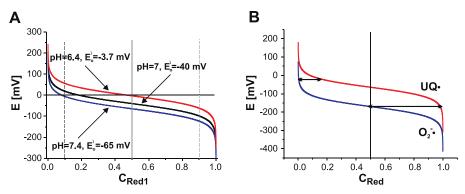


Fig. 1. Visualization of redox potential dependencies. (A) Dependence of the effective redox potential of a redox couple involving proton transfer on the portion of the normalized reduced species c_{Red1} . (B) Redox equilibrium of the pairs UQ/UQH^{\bullet} and O_2/O_2^{\bullet} . For detail see text.

of UQH• to obtain a fraction of 3×10^{-3} of produced superoxide radical. If we have only oxidized components present (e.g. O_2 and UQ) in our reaction mixture no e⁻-transfer will happen unless the superoxide or the semiquinone are formed spontaneously by other processes. It is noted, that this discussion of redox equilibrium concurs with the chemical equilibrium model in which the equilibrium constant is defined as the ratio of reactant concentrations. In this regard, the redox potential E may be viewed as a parameter that is adjustable to shift the relative concentrations of the redox couples. This perception is the basis for all voltammetric experiments, in which the potentials at electrode interfaces bias the redox conversion of reactants.

In biological processes a single electron transfer (SET) to oxygen is the first step in the formation of reactive oxygen species. Because O2 • is a rather unstable compound it undergoes additional reduction to H₂O₂, either spontaneously (via dismutation) or in a process catalysed by superoxide dismutases (SOD). H₂O₂, on the other hand, is much more stable than ${\rm O}_2$ • despite its high midpoint redox potential $E_0' \sim 1.35 \,\text{V}$ due to its slow kinetics of electron transfer, and it is a weakly oxidizing agent under physiological conditions. Its neutrality and low reactivity enable H₂O₂ molecules to cross cellular membranes, and to travel certain distances and finally to interact with particular biological targets thus acting as second messengers in many cellular processes. As for many other second messengers (including Ca²⁺), high concentrations of H₂O₂ will have toxic effects leading to cellular apoptosis and necrosis [1]. A much more deleterious ROS than H₂O₂ is the hydroxyl radical (OH•), which is one of the most reactive chemical species known. It can be formed from $H_2 O_2$ in presence of trace metals via the Fenton reaction or from $O_2^{-\bullet}$ and H_2O_2 via the Haber-Weiss reaction [2]. The very high E_0 of OH $^{\bullet}$ (2.31 V) will enable these reactive species to react with many proteins, lipids and even DNA within a living cell, finally leading to unwanted effects. Because of its high reactivity, it will immediately oxidize the molecules in its vicinity without being able to diffuse over a noticeable distance.

The basic chemical principles outlined above are universal and apply also in biological systems. Here, the situation is of course much more complex and the identification of the major sources of ROS, their localisation and the kinetics of ROS production are of major importance for the interpretation of cellular redox signaling.

2. Sources of ROS in biological systems

ROS in biological systems can be generated by the NADPH oxidase family members NOX1-5 and DUOX1 and 2, mitochondrial electron transfer chain (ETC), xantine oxidases, lipoxygenases, cyclooxygenases, heme containing proteins, such as cytochrome

P450 enzymes and also by the Fenton reaction in compartments with a high concentration of ferrous iron. The most prominent cellular sources of ROS are the NADPH oxidases and the mitochondrial ETC [1].

2.1. NADPH oxidase enzymes (NOX)

The family of NOX enzymes consists of seven members: five conventional protein complexes NOX1-5 and two dual oxidases DUOX1 and 2 [3-6]. The first characterized NOX enzyme was the NOX2 enzyme also known as phagocyte NADPH oxidase because of its high expression in granulocytes and monocyte-derived cells which use ROS for elimination of pathogens (see Fig. 4). Afterwards, other members of the NOX family were cloned and identified in many different tissues indicating a more general role for cellular ROS production. With possibly one exception (NOX4) [7], NOX enzymes are not constitutively active. They are activated upon physiological stimuli by assembly of their core NOX subunit together with accessory subunits on the phagosome- and/or the plasma membrane. This is followed by an electron transfer from cytosolic NADPH to molecular oxygen at the contralateral side leading to subsequent formation of O₂ •, which is rapidly dismutated into the more stable H_2O_2 . It is important to note that Ca^{2+} plays an important role for NOX function, not only for NOX5 and DUOX1 and 2 which contain EF hands and are directly activated by an increase in [Ca²⁺]_i [4,8,9], but also for the other members of the NOX family.

2.2. Mitochondria

While protons are pumped to the inner mitochondrial membrane (IMM) during the process of oxidative phosphorylation and ATP production, electrons are transported along the electron transfer chain (ETC) where they ultimately reduce oxygen to H₂O. The proton gradient across the IMM is essential for ATP production. Because of the relatively high difference in standard reduction potentials between complex I and complex IV of around 0.7 V, a fraction of the transferred electrons escapes and reduces molecular oxygen in their vicinity into O_2 $\, \bullet \,$. Some studies report a rather high percentage of electrons (1-2%) escaping from the ETC while other estimate a much lower fraction (<0.1%). The reason for this discrepancy may arise from differences in experimental approaches, such as use of isolated mitochondria vs. intact cells and variable external partial pressure of oxygen [2,10]. Under physiological conditions the steady state concentrations of mitochondrial O2 * which will then be dismutated by mitochondrial superoxide dismutase (mSOD) into H_2O_2 are estimated to be in low picomolar $(O_2 \cdot)$ to low nanomolar range (H_2O_2) [11].

Similar as for NOX enzymes, mitochondrial function is tightly controlled by Ca²⁺ ions [12,13]. An increase in the mitochondrial Ca²⁺ concentration leads to increased ROS production which then may affect other redox sensitive signaling pathways [12,14–17]. Further information on mitochondrial ROS production can be found in references [18–22].

3. Detection of ROS

As their name implies, ROS are very reactive and thus extremely unstable compounds. Hence, their detection in living cells is a challenging task. Presently, almost all techniques rely on detecting end-products of chemical reactions induced by ROS. Probably the most popular tools of ROS detection are fluorescent or luminescent dyes or proteins. The development and application of these sensors has been nicely described elsewhere [23–30]. In addition, ROS can be detected by electrochemistry and electron paramagnetic resonance (EPR) spectroscopy, physico-chemical techniques increasingly applied in biological studies but still not exploited to their full potential.

Here, we will briefly summarize the application of redox sensitive dyes, protein sensors and electrochemistry and describe in more detail EPR spectroscopy and its application in physiological studies.

3.1. Dyes

Fluorescent or luminescent dyes have been widely applied in biological systems for detection of ROS. Generally, they exist in an ester form such that they can be loaded in the cellular cytosol and/or organelles and indicate intracellular changes in ROS production. Some dyes, however, do not cross the plasma membrane (PM) and can be used to determine extracellular ROS concentration. A number of different dyes are available on the market, but the most popular ones are the dichlorofluorescein (DCF) based ones. Other commonly used fluorescent dyes are based on rhodamine, ethidine and the phenoxazine backbones, well known examples include Amplex Red and Amplex Ultra Red which are based on phenoxazine (resourfin). Commonly used chemiluminescent based indicators include lucigenin as a specific detector for $O_2^{-\bullet}$ while luminol has mostly been used as an indicator for general ROS production. Based on redox active substituents in the benzene ring and using the nucleophilic character of some reactive species, the development of a new generation of dyes with potentially improved characteristics has been described [31–40]. Because of the quite complex chemistry of all of these dyes, great care has to be taken in designing biological experiments and in their interpretation. For example, there is evidence that DCF may produce secondary radicals induced by its own oxidation [41].

Advantages: Relatively cheap, intracellular ROS measurement possible, detection by any kind of fluorescence or luminescence reader, can be used in any cell type

Disadvantages: Unspecific or partially specific, photosensitive, autooxidation, irreversible, interference with cellular redox state, leakage, no ratiometric measurements possible

3.2. Proteins

The invention of genetically encoded fluorescent sensors for measurements of intracellular ROS was a significant advance in the field of redox biology. Redox-sensitive, reactive cysteine containing green fluorescent proteins (roGFPs), yellow fluorescent proteins (rxYFPs) and fusions of redox sensitive oxyR with YFP (HyPer) have already been successfully used in various studies [42–47]. In addition, several Förster resonance energy transfer (FRET) based sensors

have been used as redox sensitive probes [48]. Improved versions of these sensors and creation of transgenic animals will surely bring further advances in redox signaling research. For a detailed summary of the current state of the art see [30].

Advantages: Ratiometric (HyPer and roGFP), intracellular ROS measurements, organelle targeting, relatively specific, suitable for in vivo measurements, available for further genetic engineering, transgenic animals

Disadvantages: Moderate dynamic range, pH sensitivity, interference with cellular redox state, depends on "transfectability" of cell type if not used as a knock-in construct

3.3. Electrochemistry

Electrochemical techniques have been applied to detect various reactive oxygen and nitrogen species [49–54]. Some of these utilize direct detection of diverse reactive species based on their differential redox potentials, while others use electrodes modified with molecules or proteins with electrocatalytic activity towards the species of interest. Single-cell amperometric measurements are another elegant approach for parallel detection of ROS and RNS production [55–57].

Although all of the above mentioned techniques have regularly been used in answering many biological questions, the discovery of Scanning ElectroChemical Microscopy (SECM) was a revolutionary step towards the "in-vivo" detection of ROS. SECM involves the use of a mobile ultramicroelectrode probe, with which it is possible to scan the surface of a single cell in order to get closer insight into its redox activity and surface morphology. Such experiments are performed in either an amperometric or a potentiometric mode. The topographic and chemical information about the electrochemical processes taking place at a given site of a single-cell can be obtained from the faradaic currents measured at the working ultramicroelectrode directly or after additional modification using redox mediators. Main requirements for the successful accomplishment of such electrochemical measurements are accurate placement of the tip of microelectrode(s), precise determination of the tip-to-cell distance, and the appropriate selectivity of the ultramicroelectrodes for species that may change in concentration as a result of various cellular actions [58.59].

Advantages: Possibility to detect ROS at different sites on the PM, relatively specific, detection of low ROS concentrations, fast instrumental output.

Disadvantages: Easy pollution of the working electrodes, possible interferences from other redox systems, expensive set-up, need for specialist

3.4. EPR spectroscopy

EPR spectroscopy is based on the absorption of microwave radiation by molecules with at least one unpaired electron exposed to an external magnetic field. Therefore, EPR is the method of choice to study radicals which are selectively detected in complex chemical systems prevailing in biological or physiological experiments. Such paramagnetic molecules, in the simplest case, carry a spin S = 1/2arising from a single unpaired electron which may be imagined as little dipole magnet. Hence, in a magnetic field the two possible spin states $m_S = \pm 1/2$, sometimes called "spin up" and "spin down" states, adopt different energies and populations, so that transition between the lower and the higher energy state can be induced by electromagnetic radiation providing an energy exactly bridging this energy gap (Fig. 2A). For technical reasons this absorptive process is recorded as the first derivative of an absorption line (Fig. 2C). Typically, EPR experiments are performed at magnetic fields around 350 mT (=milliTesla) and corresponding microwave frequencies of ~9.5 GHz (X-band). The latter is highly stabilized and continuously

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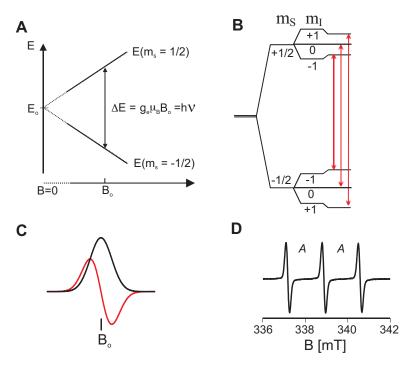


Fig. 2. Precepts of EPR. (A) The energy level separation of the two electron spin states depends on the applied magnetic field. The microwave induced transition is indicated by the arrow. (B) The electron levels are split by a 14 N-nucleus giving three EPR transitions. (C) The absorption spectrum at B_0 is recorded as first derivative. (D) The associated spectrum shows three EPR lines, A is the hyperfine splitting of 14 N.

irradiating the sample in the measuring cavity (continuous wave, cw-mode) while the field is swept in the desired range to record the spectrum.

The unpaired electron of a molecule may additionally interact with one or more nuclei in its vicinity. If these have a nuclear spin $I \neq 0$, the unpaired electron spin "is seeing" the magnetic field caused by the nuclear spins. According to the possible nuclear spin states the induced magnetic field adds to or subtracts from the applied external field, so that several lines are appearing. For a proton with I=1/2 the two spin states with $m_1=\pm 1/2$ result in four energy states for the interaction of the electron, so that two EPR lines are observed separated by the hyperfine interaction A. In many biological or physiological EPR experiments the finally observed radical spectrum is related to a nitroxide group. Here, the unpaired electron spin is located mainly in the orbitals of an NO-group and is interacting with the nuclear spin I=1 of the nitrogen. The three nuclear spin states of ¹⁴N, m_1 = +1, 0, -1, split the two electron states into a total of six energy levels which can be connected by three EPR transitions leaving the m_I-values unchanged ($\Delta m_{\rm I}$ = 0, selection rule) (Fig. 2B). This yields the characteristic three line spectrum of nitroxide based radicals (Fig. 2D). In general, from a single nucleus with spin I in an EPR spectrum 2*I* + 1 lines separated by the hyperfine constant A can be observed. When several nuclear spins are involved, the number of EPR-lines increases with the number of spin states, so that EPR spectra with many lines may be recorded. Finally, it is important that in EPR radicals can be analysed under physiological conditions at ambient temperatures or 37 °C. In aqueous solutions the recorded EPR radical spectra usually can be interpreted rather easily, because all complicating anisotropic effects of the EPR parameters are irrelevant. For measurements in solids, powders or frozen solutions a more elaborate approach to explain the EPR spectra is neces-

In principle, the radical forms of ROS (O_2^{\bullet} , HO_2^{\bullet} , OH^{\bullet}) generated in biological systems are accessible by EPR spectroscopy due to their unpaired electron spin. However, in real conditions the very

low steady state concentrations and short life times (several μs to ns) together with unfavorable relaxation properties impede their direct observation. To circumvent these restrictions, EPR spin trapping methods have been continuously developing since the mid 1960s. Up to now, an abundant variety of traps with different structures and properties have evolved for variable purposes such as chemical, biological and medicinal studies. A survey of available trapping compounds together with their application can be found in several reviews or books [2,60,61]. Their principal mode of operation is to react with the short lived radicals forming a spin adduct (or transfer a single electron), and thus adopting an unpaired electron state. This new radical or spin adduct has a considerably longer lifetime (at least in the seconds or minutes range up to several hours or days) and accumulates to well accessible concentrations easily detectable by EPR.

For measuring oxygen free radicals two major groups of spin traps are generally used, the nitrone compounds and cyclic hydroxyl amines. The chemical structures of representative spin trap reagents are shown in Fig. 3A and B. Their chemical features such as solubility, stability, trapping efficiency and cell permeability can be altered and adjusted by changing or adding specific side groups.

The central property of the nitrones (Fig. 3A) is the presence of a double bond associated with the NO group. Radicals tend to attach to this electron rich region being covalently bound to the trap structure. For example, DMPO is stabilizing the O₂ • radical at the CH-position next to NO yielding a characteristic 12-line spectrum, caused by the interaction of ¹⁴N and several nearby protons [62]. When OH• is reacting, a four-line spectrum is observed, thus allowing to distinguish the primary reactant. However, it is also observed that the DMPO-OOH adduct is transforming to DMPO-OH within minutes, so that a distinction of involved radicals is not always clear cut. The trap PBN shows a modified nitroxide three line EPR spectrum after reaction with carbon centered or OH• radicals (Fig. 3C). On the other hand, the cyclic hydroxyl amines (Fig. 3B), CPH or CMH, are transformed to a radical state by oxidation with

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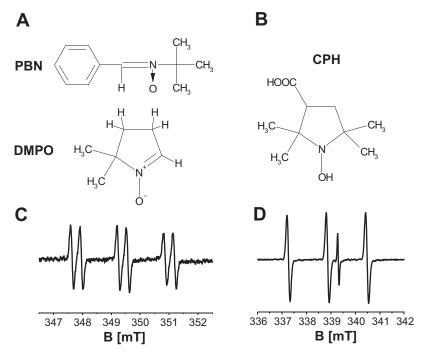


Fig. 3. Selected spin traps and their EPR spectra. Chemical structures of representative nitrone (A) and cyclic hydroxyl amine (B) spin traps together with typical EPR spectra dominated by the interaction of ¹⁴N of the NO-group of the spin adduct (cf. Fig. 2). The signals arise after trapping of carbon centered or OH*-radicals by PBN (C), or after oxidation of CPH by superoxide (D). The smaller narrow signal is from a field marker compound. CPH, 1-hydroxy-3-carboxy-2,2,5,5-tetramethylpyrrolidine; DMPO, 5,5-dimethyl-pyrroline N-oxide; PBN, N-tert-butyl-α-phenylnitrone.

 O_2 • or other oxidants like OH• yielding in all cases the same typical three line nitroxide spectrum (Fig. 3D) [63]. To distinguish what type of oxidizing species is effective, specific radical scavengers such as SOD for superoxide need to be employed. If the nitroxide signal is persistent, an oxidizing reactant like OH• or yet another species have to be considered and analysed further.

Summing up, the spin trapping technique provides a useful method to specifically identify radicals in a complex biological system via the spectral pattern of the spin adduct and/or to sensitively quantify the radical formation via the build-up of signal intensity. Ideally, a trap reagent efficiently scavenges generated primary radicals at non-toxic concentration in a biological measurement. The radical product (spin adduct) has to be sufficiently stable; it should not decay spontaneously and should be inert against further reduction or oxidation processes by other radicals or compounds (like ascorbic acid, glutathione etc.) present in the setup. In practice, the usually non-ideal properties of a spin trap in a physiological or medical experiment require well designed control experiments to avoid potential misinterpretations.

Advantages: Very selective for radicals, possibility to distinguish between different primary radicals, intra- and extracellular measurements associated with trap properties and specific radical scavengers, sensitivity in nanomolar range, easy and fast to quantify

Disadvantages: No single cell measurements possible, expensive equipment and need for specialist, short life time of spin adducts may impede radical identification and quantification

Checkpoints for controlling the intracellular ca²⁺ concentration

In virtually all cells, Ca^{2+} ions have an essential role in the regulation of many signaling cascades and cellular functions [64–66]. While some of these processes can be governed by Ca^{2+} ions in the extracellular milieu [67,68], most of them are initiated by changes in the intracellular Ca^{2+} concentration ($[Ca^{2+}]_i$). The steep (~ 10000

fold) concentration gradient between the basal $[Ca^{2+}]_i$ and the extracellular Ca^{2+} ($[Ca^{2+}]_o$), already indicates the importance of constant and tight control of cellular Ca^{2+} homeostasis. Transient increases in $[Ca^{2+}]_i$ are the essence of most Ca^{2+} -regulated cellular functions. To fully and precisely control these dynamic changes in $[Ca^{2+}]_i$, cells require a complex machinery of Ca^{2+} channels, pumps, transporters and Ca^{2+} binding proteins. These can be embedded in the plasma membrane, membranes of endoplasmic reticulum (ER), mitochondria and other organelles or are dissolved in the cytosol [64-66].

Major influx pathways for the entry of extracellular Ca²⁺ into the cytosol include voltage-gated Ca²⁺ channels (CaV) [69], receptor operated channels (such as transient receptor potential (TRP) channels [70–73] and others) and Ca²⁺ release-activated Ca²⁺ (CRAC/Orai) channels [74]. Certain TRP channels are also expressed in membranes of intracellular organelles and may thus be involved in the intracellular Ca²⁺ handling [75,76].

Because a prolonged increase in $[Ca^{2+}]_i$ is usually toxic, rapid and regulated transport of Ca^{2+} out of the cytosol is of vital importance [77]. Ion pumps located within the plasma membrane such as the Ca^{2+} ATPases (PMCA) are required for efficient Ca^{2+} extrusion. In addition, the Na^+/Ca^{2+} exchanger (NCX) and the $Na^+/Ca^{2+}-K^+$ exchanger (NCKX) also play an important role. Under certain conditions these exchangers can reverse their mode and contribute to an increase of $[Ca^{2+}]_i$ [64–66]. Besides extruding Ca^{2+} to the extracellular space, intracellular Ca^{2+} stores also play an important role in regulating Ca^{2+} homeostasis. The most prominent stores are the lumen of the ER and the mitochondria.

Within its lumen, ER stores contain Ca^{2+} at concentration of \sim 0.5 mM, which can be rapidly released into the cytosol through activated inositol 3-phosphate-(IP₃R) or ryanodine receptors (RyR) [78–80]. The refilling of these Ca^{2+} pools is achieved by sarcoendoplasmic reticulum Ca^{2+} ATPases (SERCA). SERCAs pump Ca^{2+} back into the ER, as some Ca^{2+} constantly leaks across the ER membrane [81–85]. The molecular identity of these leak channels is still

controversial, although recent data points towards the involvement of the Sec61 translocon [86–88].

In mitochondria, Ca²⁺ ions cycle between the cellular cytosol and the mitochondrial matrix thereby regulating organelle function and ATP production [89–91]. During this process Ca²⁺ ions have to cross two membranes: the outer- (OMM) and the inner mitochondrial membrane (IMM). It has been proposed that the OMM is porous and therefore does not play a major role in mitochondrial Ca²⁺ entry. However, some studies suggested that the voltage dependent anion channel (VDAC) might play an important indirect role in the mitochondrial Ca²⁺ homeostasis [92–94]. Across the IMM, two major transport pathways for Ca²⁺ ions are proposed. One is activated at relatively high intramembrane space Ca²⁺ concentration and is underlined by a very selective channel known as the mitochondrial Ca²⁺ uniporter (MCU). The second pathway has much lower capacity but higher Ca²⁺ sensitivity and is probably governed by several channels or transporters [95].

Very recently, two groups identified an IMM-based, two-transmembrane domains containing protein known as CCDC109A as a pore forming unit of MCU, which is a great advance in the field of mitochondrial Ca²⁺ regulation. However, the orientation of the protein in the membrane and its role either as a bona fide ion channel or as part of a larger protein complex still needs to be clarified [96,97]. In addition, an IMM protein known as Letm1 has been shown to act as an efficient Ca²⁺/H+ exchanger leading to an increase in the Ca²⁺ concentration or the mitochondrial matrix [98]. We have also shown that hydroxylated coenzyme Q (CoQ) forms (present at the IMM) are able to transport Ca²⁺ across artificial

membranes but if and how these CoQs affect the mitochondrial Ca^{2+} homeostasis is not yet clear [99,100]. Given the biophysical properties of Ca^{2+} transport by Letm1 and hydroxyl CoQs, it is possible that they are involved in the slow mitochondrial Ca^{2+} uptake.

To remove Ca^{2+} from the mitochondrial matrix and prevent Ca^{2+} overload and subsequent pathological conditions the Na^+/Ca^{2+} exchanger is essential [90]. Similar as for Ca^{2+} import, some reports suggest a role for the mitochondrial permeability transition pore (mPTP) and the OMM for mitochondrial Ca^{2+} efflux [101].

Obviously, many of the Ca²⁺ transport and binding processes are now understood on the molecular level. This is one prerequisite to understand their interplay to control the Ca²⁺ homeostasis during cell activation. ROS modulate many transporters and are therefore involved in the modulation of Ca²⁺ homeostasis (Fig. 4).

5. Effects of ROS on ion channels

5.1. General mechanisms of protein oxidation

While ROS can react with nuclear DNA, lipids of the cellular membranes and with polypeptides, it is mainly their action on proteins that endows them with specificity. The wide range of cellular functions that are regulated by ROS signaling include – among others – growth factor responses, cell adhesion and apoptosis, and depend on specific and reversible modification of the proteins involved in the related signaling cascades [2]. A

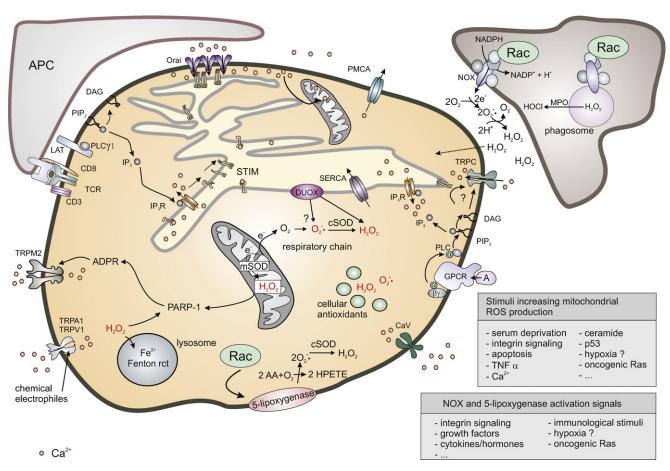


Fig. 4. Redox and Ca²⁺ signaling pathways. A, agonist; APC, antigen presenting cell; CD, cluster of differentiation; DAG, diacylglycerol; DUOX, dual oxidase; GPCR, G protein-coupled receptor; IP₃, inositol 1,4,5-trisphosphate; IP₃R, IP₃ receptor; LAT, linker of activated T cells; MPO, myeloperoxidase; PARP-1, poly-(ADP-ribose)-polymerase; PIP₂, phosphatidylinositol 4,5-bisphosphate; PLC, phospholipase C; PMCA, plasma membrane calcium ATPase; SERCA, sarco-endoplasmic reticulum ATPase; STIM, stromal interaction molecule; TRP, transient receptor potential; cSOD, cytoplasmic superoxide dismutase; mSOD, mitochondrial superoxide dismutase; ADPR, ADP-ribose; AA, arachidonic acid; HPETE, hydroperoxyeicosatetraenoic acid; VOC channel, voltage operated calcium channel; NOX, NADPH oxidase; NADP, nicotinamide adenine dinucleotide phosphate.

well-studied example is the response of E. coli to oxidative stress: the oxidation of two–SH groups by increasing concentrations of H_2O_2 leads to reversible formation of a disulphide bond within the transcription factor oxyR, thereby activating the expression of several antioxidative enzymes, which protect E. coli against increased levels of oxidative stress [102,103].

High concentrations of ROS can cause a wide range of amino acid modifications, e.g. the oxidation of thiol groups to sulfenic, sulfinic and sulfonic acids, the oxidation of arginine and lysine residues to aldehydes or of methionine to the corresponding sulphoxide or sulphone. Oxidation can also lead to intra- and intermolecular crosslinking and trapping of proteins in multimolecular complexes. Many of those modifications are essentially irreversible, abolish protein function and eventually lead to degradation. On the contrary, physiologically relevant concentrations of ROS can react with specific regulatory amino acids within the polypeptide chain. This selective action depends on the accessibility within the tertiary structure, the local concentration and nature of the reacting redox species and on the local redox potential. While redox signaling regulates the function of an array of proteins ranging from transcription factors to protein kinases and phosphatases [2,14,104–106], we will focus here on Ca²⁺ conducting ion channels, in particular voltage-gated Ca²⁺ channels, the TRP superfamily and CRAC/Orai channels.

5.2. Voltage-gated Ca²⁺ channels

The voltage-gated Ca^{2+} channel (CaV) family consists of 5 major subgroups known as L-type- (Cav1.1–1.4), N-type- (Cav2.2), P/Q-type- (Cav2.1), R-type- (Cav2.3), and T-type- (Cav3.1–3.3) Ca^{2+} channels. They are expressed in many cell types and are responsible for various cellular functions such as muscle contraction, control of action potential, secretion, gene expression and others. Because both redox- and Ca^{2+} -signaling play an important role in the cardiovascular system, where CaV channels are highly expressed, they were one of the first Ca^{2+} channels to be identified as redox sensitive.

In a number of studies a large variety of oxidants have been found to regulate CaV channel activity on different levels, affecting their expression, trafficking, open time and open probability. Some of these studies could demonstrate that reactive cysteines in the pore-forming $\alpha 1$ -subunit are the molecular targets for ROS. CaV channel redox regulation has been studied in myocytes and neurons of various species as well as in heterologous expression systems. Similar as for the reports on redox regulation of Orai as well as for some TRP channels (see below), some authors described activation of CaV by oxidation while others reported the contrary, i.e. inhibition of channel activity.

These differences may depend on the cell type and genetic modulation (endogenous channels or an over-expressing system) but also point towards a complex redox regulation of Ca²⁺ signaling in the cardiovascular and nervous system where oxidation of additional targets could influence CaV channel function.

The findings regarding the redox regulation of CaV channels are summarized in Table 1 and have been described elsewhere in detail [107–109].

5.3. TRP channels

Members of the mammalian transient receptor potential (TRP) family of ion channels are subdivided into seven subgroups according to sequence homologies and protein domains (TRPC, TRPM, TRPV, TRPA, TRPN, TRPP and TRPML). The canonical or "classical" TRPC members share the highest degree of sequence homology to the founding member *Drosophila* TRP. TRP channels are cation-permeable and are involved in the detection of differ-

ent sensory modalities. Their activating mechanisms include direct detection of temperature and/or chemical compounds, extra/intracellular ion concentrations but also detection of second messengers generated by G-protein coupled receptors (GPCR) cascades. The expression pattern of TRP channels range from ubiquitously expressed (i.e. TRPM7) to very restricted expression in specialized cells [71,110,111]. While all cells can be exposed to intracellular ROS, exposure to extracellular ROS generated by environmental factors such as UV light or chemical irritants or by specialized ROS-producing cells, such as monocyte derived cells, granulocytes and thyroid cells depends on localization. Cells and tissues in the immediate vicinity of the ROS source will obviously be affected more than others.

Regarding regulation of ion channels by oxidation, TRPM2 channels have been most extensively investigated (see below and Table 2), however, other TRP channels such as TRPV1 and TRPA1, expressed in sensory neurons mediating responses to irritant substances such as pungent and electrophilic compounds found in chili peppers, mustard oil, onions, garlic and cinnamon extracts [112–117], are also excellent targets for investigation of cysteine modifications. Indeed, pain related behavior relayed by TRPA1 in wildtype but not in TRPA1 -/- mice can be aggravated by irritant, oxidizing compounds such as allicin and diallyl disulfide (DADS), but also by H₂S and H₂O₂ [114,118,119]. In an effort to identify the targets of the electrophilic attack in heterologous expression systems, two independent groups used cysteine mutagenesis studies. Hinman et al. [120] identified three critical cysteine residues (C621, C641, and C665*) downstream of the ankyrin repeat region of human TRPA1 as being critical in conferring sensitivity towards Nmethylmaleimide (NMM), although the mutant retained a residual sensitivity towards allyl isothiocyanate (AITC), which was annulled by further mutation of K710 [120], (*note: the original report has a-2 shift in the amino acid nomenclature). In a parallel study, Macpherson et al. [121] used click chemistry and mass spectrometry to identify 14 cysteines that can be modified by iodoacetamide (IA); three of which (C415, C422 and C622, mouse TRPA1) are critical for mediating the activating effect of mustard oil on channel function. Cysteine 622 (mTRPA1) is identical to C621 (hTRPA1) and is conserved in TRPA1 clade members with the exception of Ciona intestinales, as also C622, C415, C422, C642 and K712 (mTRPA1), suggesting an ancient shared origin of chemical nociception [122].

In the case of the capsaicin receptor TRPV1, allicin, one of the active compounds in garlic and onions, was found to sensitize TRPV1. In a thorough analysis of the actions of allicin on TRPV1, Salazar et al. [123] found that, in contrast to TRPA1 channels, modification of only a single cysteine (C157) located in the N-terminal region of TRPV1 was necessary and sufficient to mediate the activating effect of allicin or MTSEA. Interestingly, in both TRPA1 and TRPV1 the modified cysteines are localized in cytosolic regions of the proteins, thus require the electrophilic compounds to cross the plasma membrane. Besides the likely physiological role of these cysteine modifications in the sensation of the burning feeling of spicy foods, oxidant (reactive electrophiles) induced modification of TRPA1 residues plays an important role in the chemical nociception of toxic compounds (e.g. during cigarette smoking) and represents an evolutionary ancient detection system [122]. TRPA1 is thus also an attractive candidate for the development of drugs to reduce pain and inflammation associated with airway inflammation.

5.3.1. Redox activation of TRP channels in cell death

The best studied target for ROS induced cell death in mammals is TRPM2, a ubiquitously expressed member of the TRPM family. A very recent and detailed review describes the role of TRPM2 in oxidative stress [124], thus we will only provide a brief summary here. TRPM2 is a non-selective cation channel with a permeability of Na⁺ > Ca²⁺ > K⁺ and is activated by cADPR, ADPR and

 Table 1

 Redox regulation of voltage gated calcium channels.

	Cells	Redox modulator	Effect	Target	Ref.
L-type	HEK, rat myocytes guinea pig, myocytes, frog myocytes, PC12, neurons	PHMPS, thimerosal, 2.3 mM purine + 20 U/I XO, DHF, ascorbate/Fe ²⁺ , PCMBS, MTSEA, DTNB, X/XO, Endogenous ROS, Hypoxia, H ₂ O ₂	↑ Ca ²⁺ currents	cysteine oxidation, increased trafficking, upregulation of channel expression	[219–223]
	HEK, CHO, rat myocytes, guinea pig myocytes, arterial myocytes, chick retinal cells	H ₂ O ₂ , MTSEA, MTSET, PHMPS, DTDP, DTBNP, 2,2'-dithiodipyridine thimerosal, SIN-1, DTNB, H ₂ O ₂ /Fe ³⁺ , hypoxia, CO, endogenous ROS, CORM-2	↓ Ca ²⁺ currents	cysteine oxidation, C1789, C1790, C1810 on α1C, reduction in dihydropyridine binding sites, reduced sensitivity towards physiological stimulation, changes in PM potential	[224-232]
T-type	HEK, rat neurons HEK, rat neurons, rat peripheral nociceptors	GSH, DTNB, L-cysteine, DTT, TCEP GSSG, MTSET	↑ Ca ²⁺ currents ↓ Ca ²⁺ currents	extracellular cysteines	[233–237]
P/Q-type	xenopus oocytes	${ m H_2O_2},{ m DTNB}$	↓ Ca ²⁺ currents ↑ Ca ²⁺ currents at positive potentials	ND	[238]

CHO, chinese hamster ovary; CORM-2, tricarbonyl dichloro ruthenium(II) dimer (CO donor); DHF, dihydroxyfumaric acid; DTBNP, 2,2'-dithiobis(5-nitropyridine); DTDP, 2,2'-dithiodipyridine; DTNB, 5,5'-dithiobis[2-nitrobenzoic acid]; DTT, dithiothreitol; GSH, glutathione; GSSG, glutathione disulfide; HEK, human embryonic kidney; MTSEA, 2-aminoethyl methanethiosulfonate hydrobromide; MTSES, sodium (2-sulfonatoethyl) methanethiosulfonate; MTSET, [2-(Trimethylammonium)ethyl] methanethiosulfonate bromide; PCMBS, *p*-chloromercuribenzene sulphonic acid; PHMPS, *p*-hydroxy-mercuric-phenylsulphonic acid; SIN-1, linsidomine (NO donor); TCEP, tris(2-carboxyethyl)phosphine; X, xanthine; XO, xanthine oxidase.

NAADP [73,125–127]. Mori and coworkers first described activation by addition of H_2O_2 , leading to cell death [128]. Interestingly, ROS mediated activation of TRPM2 is not a direct effect, but is mediated by an increased mitochondrial synthesis of the TRPM2 ligand ADP-Ribose [129] as well as an increased ADP-ribose synthesis by the nuclear enzyme poly-(ADP-ribose)-polymerase PARP-1, no oxidant-induced activation of TRPM2 is detected in PARP-1 deficient cells [130]. In monocytes, which are able to produce significant amounts of H_2O_2 through NOX2, activation of TRPM2 triggers an Erk mediated pathway that ultimately leads to enhanced transcription of CXCL8, a chemokine that leads to neutrophil infiltration at sites of inflammation. TRPM2 knock-out mice are protected against this infiltration and show reduced inflammation in a mouse model of colitis ulcerosa [131]. Neurons also express TRPM2 where it may be involved in H_2O_2 -induced neuronal death [132,133].

In contrast to oxidant-induced activation of the Ca^{2+} permeable TRPM2 which leads to apoptosis, a recent report describes redox dependent facilitation of TRPM4 currents by H_2O_2 which can ultimately lead to cell death by necrosis. Down regulation of TRPM4 in HeLa cells prevented H_2O_2 induced necrotic cell death while leaving apoptotic pathways intact [134]. In contrast to all other TRP channels, TRPM4 and TRPM5 are monovalent-selective cation channels that are Ca^{2+} activated and likely regulate membrane potential [135,136]. Because both TRPM2 and TRPM4 are activated or facilitated in a similar concentration range of H_2O_2 ($EC_{50} \sim 50 \,\mu\text{M}$), relative expression levels of these channels, of PARP-1 and of intracellular antioxidant proteins might determine whether and when cells undergo oxidant induced apoptosis versus necrosis. While TRPM2 oxidant induced activation is indirect, hTRPM4 is directly modulated by oxidation of a C-terminally

Table 2Redox regulation of TRPM, TRPA and TRPV channels.

Channel	Cells	Redox modulator	Effect	Target	Ref.
TRPM2	HEK, neurons, microglia, A172, vascular endothelial cells	H ₂ O ₂ 0.1-5 mM, nitrosothiol donor MNNG 10 μM	↑ Ca ²⁺ , Na ⁺ influx, currents ↑ apoptotic cell death	no direct target, increased production of ADPR (mitochondria)	[128–130], [132] [239–243]
TRPM4	нек, нега	H ₂ O ₂ 0.2 mM	↑ Ca ²⁺ , Na ⁺ influx desensitization ↑ necrotic cell death	direct, target intracellular C1093	[134]
TRPM6	НЕК	H ₂ O ₂ 1 mM	↓ outward current	direct, blocked by intracell. interaction with MsrB1 M1755	[244]
TRPM7	HEK, neurons	OGD, NaCN, H ₂ O ₂ , menadione, NO	↑ inward Ca ²⁺ , Na ⁺ current ↓ outward current	no direct effect, activity of TRPM7 increases ROS production	[143]
TRPA1	HEK, C-neurons WT/KO mice bladder	$H_2O_2~(EC_{50}\sim 230~\mu M),$ AITC, NMM, mustard oil, MTSEA, MTSET, AI, CA. NaHS	↑ Ca ²⁺ , Na ⁺ current	direct, intracellular C415, C422, C622 C619, C639, C663, K708	[117–121] [245–247]
TRPV1	sensory neurons, WT/KO mice, HEK, Het1A	thimerosal, H ₂ O ₂ 1 mM	↓ Ca ²⁺ , Na ⁺ current	extracellular: C621 G.g. specific: C66, C111, C832	[123,246,248,249
		H_2O_2 1 mM, allicin, MTSEA, HNE	↑ Ca ²⁺ , Na ⁺ current	direct intracellular C157, G.g.: C783	[248,250]

A172, human glioblastoma cell line; AI, allyl isothiocyanate; AITC, allyl isothiocyanate; CA, cinnamaldehyde; HEK, human embryonic kidney; HeLa, human andenocarcinoma cell line; Het1A, human epithelial cell line (esophagus); HNE, 4-hydroxy-2-nonenal; MNNG, N-Methyl-N'-nitrosoguanidine; MTSEA, 2-aminoethyl methanethiosulfonate hydrobromide; MTSET, [2-(Trimethylammonium)ethyl] methanethiosulfonate bromide; NMM, N-methylmaleimide; OGD, oxygen glucose deprivation

located cysteine, C1093 (see Table 1). This cysteine residue is conserved in TRPM3, 6 and 7, although its role has not been investigated within these channels.

TRPM6 and TRPM7 are both unusual members of the TRP family as they possess both ion channel and protein kinase activities. These serine/threonine kinase domains are located at the extreme C terminus, but their activity does not directly affect channel function, although autophosphorylation has been demonstrated for TRPM7 [137]. Both channels are divalent selective, allowing entry of Mg²⁺ and Ca²⁺, but will be inhibited by intracellular Mg²⁺ (0.3–1 mM) [138,139]. While TRPM7 is ubiquitously expressed and essential for Mg²⁺ homeostasis and cell survival, TRPM6 is strongly expressed in kidney and intestine where it also plays an important role in epithelial Mg²⁺ reabsorption [140,141]. TRPM6 interacts through its alpha-kinase domain with methionine sulfoxide reductase B1 (MsrB1). This protein interaction is able to mask the inhibitory effect of H₂O₂ on channel activity. Mutation of methionine 1755 within the kinase domain was able to reduce the inhibitory effect of H₂O₂, thereby mimicking the action of MsrB1. Reversible oxidation of methionine residues is a relatively novel concept in ion channel physiology. Methionine oxidation has been reported to disrupt inactivation of shaker voltage-dependent K⁺ channel activity, an effect that can be reversed upon coexpression with MsrA1 [142]. Whether a similar mechanism can occur in TRPM7 is unknown. On the contrary, suppression of TRPM7 expression reduced cell death induced by prolonged oxygen glucose deprivation (OGD) in neuronal cultures. OGD activated Ca²⁺ permeable non-selective currents that resemble those of heterologously expressed TRPM7. The authors demonstrated enhanced inward currents in H₂O₂ (1 mM) stimulated TRPM7 expressing HEK cells [143]. A finding that several independent siRNA approaches to reduce TRPM7 in neurons also decreased expression of TRPM2 mRNA, raised the possibility that TRPM7 and TRPM2 expression are interdependent, thus further complicating the analysis of the effects of oxidative stress on native cells. A summary of redox regulation of TRPM2, -4, -6, -7, A1 and V1 channels is given in Table 2

5.3.2. Redox regulation of TRPC channels

We will only briefly summarize the effects of ROS on TRPC channels as several reviews recently summarized the existing data [144-147]. One of the earliest reports on redox regulation of TRP channels by the Groschner group reports regulation of TRPC3 by the oxidant tBHP (tert-butylhydroperoxide). tBHP increases cation currents in porcine aortic endothelial cells, an effect that has also been observed in cells overexpressing TRPC3 or TRPC4 channels [148]. The subsequent finding that the activating effect of oxidants in TRPC3 expressing cells can be inhibited by the U73122, an inhibitor of phospholipase C (PLC) [149], might suggest that activation does not involve direct oxidation of thiols in TRPC proteins, although a required costimulatory role of phospholipids such as diacylglycerol (DAG) together with a direct oxidant effect cannot be ruled out. Because ROS are known to inhibit protein phosphatases such as PP2A, the effects of increased ion channel phosphorylation versus a direct oxidation of ion channel thiol-groups need to be taken into account.

A further complication in the analysis of reactive cysteines results from the cross-talk between the oxidation of cysteines by ROS and cysteine S-nitrosylation by nitric oxide. Mild (reversible) oxidation as well as rapid and highly instable formation of S-nitrosylated cysteines by NO which can be generated by hypoxic conditions or through activation of cellular nitric oxide synthase (NOS) may be viewed as a posttranslational modification such as phosphorylation. In a comprehensive investigation the role of nitric oxide (NO) and H₂O₂ on TRP channels was analysed [150]: The receptor operated TRP channels (TRPC5 and C7) as well as the thermo-sensitive TRPV channels (TRPV1, V3 and V4) all increased

Ca²⁺ entry into cells in response to the NO donor SNAP, an effect which is partially reversed by additional application of the reducing agent DTT. The effect is mediated by cysteine S-nitrosylation of the conserved cytoplasmatically accessible C553 and C558 of the H7 pore loop region between S5 and S6 of TRPC5 likely bending the S6 segment to open the intracellular activation gate. The membranepermeable thiol-reactive compound MTSEA can mimic this effect, but not the membrane-impermeable MTSET. Similar effects of NO can be seen for TRPV1, TRPV3 and TRPV4 that contain cysteines within the same region. In addition, Yoshida et al. showed that TRPC5 is activated by H₂O₂ as well [150]. Another study investigated the effects of the intracellular and secreted redox active protein thioredoxin on TRPC5 channel function. The authors report that extracellular application of reduced thioredoxin as well as of DTT can activate TRPC5 by breaking an existing disulfide bridge between C553 and C558 [151], which generates a discrepancy as to how accessible these cysteines in the S5-S6 linker region are towards either the intra-or extracellular side. The combined groups of Yao and Beech were unable to reproduce the H₂O₂-mediated activation of TRPC5 [152]. However, in a recent publication, the Beech group did indeed show TRPC5 activation by H₂O₂, an effect that was suppressed by exogenous anti-oxidants [153]. Considering the results discussed above, it would be interesting to investigate if the pore loop exists as an unstable equilibrium between reduced (redox sensitive) C553 and C558 and a disulfide bonded C553 with C558, thus allowing the pore to adjust to different oxidative environments, which depending on culture and recording conditions, may yield diverging results as to their redox sensitivity. A summary of redox regulation of TRPC channels is given in Table 3.

5.4. Store operated Ca²⁺ entry (SOCE) and CRAC/Orai channels

The concept of store operated Ca²⁺ entry (SOCE) was first proposed by Jim Putney in the mid eighties [154,155]. Soon afterwards, several seminal papers described signaling events underlying SOCE and the biophysical properties of the Ca²⁺ channel itself. The channel was named Ca²⁺ release activated Ca²⁺ (CRAC) channel with the ion current described as (I_{CRAC}) [156–158]. The molecular identity of the CRAC channels was not known until 2006 when three independent groups identified the pore forming protein [159-161]. Besides the major component Orai1 (CRACM1), which when mutated (R91W) leads to severe combined immune deficiency (SCID) [159], two homologs, namely Orai2 and Orai3 have been identified. Their physiological function is much less defined. Orai homologs can be activated by the Ca²⁺ sensing ER-proteins STIM1 and STIM2, which were already identified as necessary for activation of SOCE [162-164]. The signaling cascade underlying SOCE thus spans from receptor mediated activation of PLC, hydrolysis of phosphatidylinositol 4,5-bisphosphate with subsequent IP₃ production, IP₃R mediated depletion of ER Ca²⁺ stores which in turn leads to a conformational change, oligomerization and translocation of STIM1 to plasma membrane near regions of the ER where STIM1 activates Orai channels through direct interaction with its C-terminal region (Fig. 4). A number of groups have also reported the participation of TRPC1 and TRPC3 in a STIM mediated storeoperated Ca²⁺ entry pathway, so caution is needed when analyzing native SOCE in the absence of the clearly identified molecular components [165,166].

Redox modification of any of these components may ultimately result in alterations in SOCE. Moreover, STIM1-induced modulation of TRP or voltage-gated Ca²⁺ channel [167,168] activity may further complicate analyses of ROS mediated effects. Also to be taken into consideration are indirect effects such as ROS mediated activation of cation channels which depolarize the PM and thus lead to indirect modulation of CRAC channel activity by reducing the driving force for Ca²⁺ influx or the influence of altered phosphorylation of SOCE

Table 3Redox regulation of TRPC channels.

Channel	Cells	Redox modulator	Effect	Target	Ref.
TRPC3	ECAP	tBHP 0.4 mM	↑ Na ⁺ influx	ND, blocked by La ³⁺	[148]
TRPC5TRPC5_C1(TRPC7TRPM2)	HEK, endothelial cells, VSMC, BAEC	SNAP 0.3 mM, $\rm H_2O_2$ 1 mM, oxidized phospholipids, thioredoxin, DTT 10 mM	↑ Na ⁺ , Ca ²⁺ influx ↑ Na ⁺ , Ca ²⁺ influx upon disulfide bond breakage	direct, S-Nitrosyl. C553, C558 direct target S-S bond C553-C558	[150–153]
TRPC6	HEK, kidney glomeruli	PAN (NOX activator), H_2O_2 EC $_{50}\sim50~\mu\text{M},$ NEM, thimerosal	↑ Na ⁺ , Ca ²⁺ influx ↑ expression	ND indirect, no effect on single channel	[251,252]

BAEC, bovine aortic endothelial cells; DTT, dithiothreitol; ECAP, porcine aortic endothelial cells; HEK, human embryonic kidney; NEM, *N*-ethylmaleimide; PAN, puromycin aminonucleoside; SNAP, *S*-nitroso-*N*-acetyl-DL-penicillamine (NO donor); tBHP, *tert*-butylhydroperoxide; VSMC, vascular smooth muscle cells.

proteins. The activity of Ca²⁺ pumps (PMCA) are also affected by oxidation, although at higher concentrations of oxidants and also depending on the cell type [169,170].

5.4.1. Reactive oxygen species effects on SOCE and CRAC/ORAI channels

The first indications that redox processes could affect SOCE were reported when thiol group modifying agents, such as thimerosal, were shown to sensitize the IP₃R but also induce I_{CRAC} independently of IP₃ [171–174]. These findings were followed by several studies leading to the conclusion that oxidation of IP₃R and RyR induces their activation and thus lead to depletion of the ER Ca²⁺ stores (for more details see [107,108] and Table 4). On the other hand, there are few studies reporting redox regulation of SOCE independently of IP₃R or RyR. Elliot et al. [175] describe a complex and time dependent effect of the oxidizing agent tBHP (0.4 mM) on Ca²⁺ signaling in endothelial cells. They show that short incubation with tBHP (less than 1 h) neither affected the induced ER Ca²⁺ store content nor its depletion but suppressed SOCE. Longer incubation times (1–2 h) reduced store depletion in addition to causing SOCE inhibition, whereas more than 2 h tBHP incubation induced higher basal [Ca²⁺]_i and also a prolonged Ca²⁺-signal following SOCE activation. Several years later Törnquist et al. confirmed the finding that SOCE is inhibited by thiol-modifying oxidizing agents such as thimerosal, tBHP and H_2O_2 (300 μ M). However, their results suggest that these effects are mediated through activation of protein kinase C (PKC) [176,177]. Vanadate and its derivative pervanadate obtained from a chemical reaction with H2O2 are well established as thiol modifying agents and also implicated in modulating SOCE. However, these compounds are also very potent phosphatase inhibitors and oxidants could thus regulate SOCE via increased phosphorylation. Indeed, several studies showed that maintenance of SOCE can be modulated by redox induced changes in phosphorylation [14,178,179]. On the other hand, Ehring et al. reported that vanadate compounds act in a similar way as thimerosal: They deplete the ER Ca²+ stores via thiol oxidation and thus induce activation of I_{CRAC} independently of any kinases or phosphatases [180]. Additional evidence that SOCE is inhibited by oxidizing agents such as homocysteine, H_2O_2 , HOCl, tBHP and diamide was later provided by several independent groups in different cell types [181–184]. Particularly interesting are the findings by Redondo et al. who observed modest activation of SOCE following treatment with low μM concentrations of H_2O_2 while higher concentrations (low mM) inhibited SOCE in human platelets [185]. Furthermore, Suzuki et al. observed activation of SOCE by O_2 • but suppression by H_2O_2 in mast cells [186].

However, the molecular mechanism(s) underlying the different effects of oxidation and thiol modification on SOCE remained relatively unclear. One of the major reasons for this, as mentioned above, was the unknown molecular identity of the CRAC channels. Recently, using electrophysiology, Ca²⁺-imaging and site-directed mutagenesis we found that Orai1, but not Orai3 activation is inhibited by preincubation with H₂O₂ [187]. Cysteine 195 which is present in Orai1 but absent in Orai3 was identified as a major redox sensor of Orai1, although C126 and C143 also contributed to redox sensitivity. Mutation of all three cysteines in Orai1 render the channels functional but completely insensitive to preincubation with H_2O_2 , providing evidence that in a heterologous expression system, longer extracellular preincubation with H₂O₂ mediates its effects on CRAC solely through oxidation of Orai1. Increased expression of the insensitive Orai3 enables primary effector T cells to reduce their SOCE sensitivity towards extracellular H₂O₂ [187]. Acute application of H₂O₂ to T cells also induced a rise in internal Ca²⁺, which is likely independent of Orai1, as siRNA knock-down of Orai1 did not reduce this acute increase in intracellular Ca²⁺ (unpublished data).

Hawkins et al. [188] studied redox mediated activation of STIM1 and found that oxidative stress leads to gluthationylation of STIM1's cysteine 56, triggering STIM1 oligomerization and punctae

Table 4Redox regulation of SOCE and CRAC/Orai channels.

SOCE	Cells	Redox modulator	Effect	Target	Ref.
Orai1	HEK, CD4⁺ T cells, Jurkat T cells	H_2O_2 $EC_{50}\sim 34$ and $40\mu M$	↓ Ca ²⁺ influx ↓ Ca ²⁺ current	C126, C143, C195	[187]
STIM1	HEK, smooth muscle, neurons, COS7, DT40 B-cells, MEF, 143B osteosarcoma	H ₂ O ₂ , hypoxia, BSO	↑ Ca ²⁺ influx ↑ Ca ²⁺ current	C56	[188,200–202]
IP ₃ R, RyR unidentified target	RBL, HEK, Jurkat, endothelial cells, FRTL-5, SH-SY5Y, platelets, smooth muscle, neutrophils, mast cells, DT40 B-cells	H ₂ O ₂ , hypoxia, thimerosal, tBHP, vanadate, O ₂ -•, pervanadate, HOCl, NO, DEANO, GEA3162, SNP, NO ₂ -	$\uparrow\downarrow$ Ca ²⁺ influx $\uparrow\downarrow$ Ca ²⁺ current (see text for details)	IP₃R and RyR reactive cysteines and ND	[171–177,180–186,189, 192–195,203–218]

BSO, buthionine sulfoximine; DEANO, 2-(*N*,*N*-diethylamino)diazenolate-2-oxide; FRTL-5, fischer rat thyroid low serum 5%; GEA3162, 1,2,3,4,-oxatriazolium, 5-amino-3-(3,4-dichlorophenyl)-chloride; HEK, human embryonic kidney; IP₃R, IP₃ receptor; Jurkat, human leukemic T cell line; MEF, mouse embryonic fibroblasts; RBL, rat basophilic leukemia; RyR, ryanodine receptor; SH-SY5Y, human neuroblastoma cell line; SNP, sodium nitroprusside; tBHP, *tert*-butylhydroperoxide.

formation independently of the ER Ca^{2+} concentration. Clustered STIM1 proteins are then able to activate Orai channels and induce Ca^{2+} entry across the PM. STIM1 C47/C56 mutants are constitutively active, although these effects may be due to defective STIM1 EF hand function. Another study by Grupe et al. reports that H_2O_2 directly mediates store depletion and thereby activates Orai in a STIM1 dependent manner [189], a finding that we could not reproduce in a heterologous expression system using patch-clamp analyses.

How might these discrepancies be resolved? A number of recent studies investigated the influence of the STIM1:Orai stoichiometry [190,191] on current size, which implies that different STIM:Orai ratios may show differences in their regulatory behavior. It is also of importance to find out under exactly which conditions cysteines in STIM1 and/or STIM2 are oxidized or reduced and what kind of chemical modifications they undergo (e.g. nitrosylation or glutathionylation). In addition, STIM may activate other channels (i.e. TRPCs) leading to an increase in intracellular Ca²⁺, although their current-voltage relationship should differ from those mediated by Orai (see also Table 4 for a summary of redox regulated activation/inhibition of SOCE).

Another possible explanation of the divergent findings might be that autocrine, intracellularly produced oxidants may favor depletion of the ER Ca²⁺ stores and STIM oligomerization leading to activation of the CRAC channels, whereas paracrine extracellular production of ROS would first target Orai1 channels (C195) and hence inhibit their activation.

Further precise and detailed studies are needed to shed more light on these complex signaling events. It will also be of importance to clearly define and quantify the types of ROS using the methods outlined above. Application of one kind of ROS in a particular concentration may also change the amount of other ROS, complicating the interpretation of data.

5.4.2. Hypoxia effects on SOCE and CRAC/ORAI channels

Oxygen deprivation is a condition of immense clinical importance and a large number of studies investigate the effects of hypoxia on signaling cascades. Here, we will only focus on those aspects of hypoxia that relate to SOCE. Several studies report modulation of SOCE under hypoxic conditions in different cell types (Table 4). While many studies do not investigate the link between hypoxia and the redox state, some indicate that the effects of hypoxia occur via changes in the cellular ROS production. Currently, there is no consensus about the concentrations of ROS induced by hypoxia.

Inhibition of SOCE in hypoxic conditions was shown in several studies [192–195]. Interestingly, two of these studies used chronic hypoxia rat models and not acute induced hypoxia of cells in culture. The molecular mechanism of inhibition was however not identified. On the other hand, Asai et al. showed that SOCE is inhibited by hypoxia through extracellular, but not intracellular acidification [194].

Activation of SOCE by hypoxia was mostly reported upon acute exposure of cultured cells to low amounts of oxygen (2–4%). Two studies suggested that this activation is governed by up-regulation of TRP channels [196,197] while others localized the ER as the target-site [198–202].

Lu et al. proposed that SOCE activation by hypoxia is via STIM1 while STIM2 played no significant role in this context [200]. Conversely, Berna-Erro et al. [201] reported that STIM2 is essential for the hypoxia induced Ca²⁺ overload, while manipulations of STIM1 expression had no significant effect. One possible explanation for this discrepancy may be the different cell types used. Lu et al. performed their study in arterial smooth muscle cells while Berna-Erro et al. worked with neurons.

5.4.3. Reactive nitrogen species effects on SOCE and CRAC/ORAI channels

The focus of this review is on the role of ROS as modulators of ion channels function. Nevertheless, reactive nitrogen species (RNS) have also been implicated as modulators of ion channel function and their effects on SOCE should not be neglected. Because the targets (reactive cysteines) of ROS and RNS are often identical we will also shortly summarize the present knowledge on the impact of RNS on SOCE. Similar to the effects of ROS and hypoxia, some reports describe inhibition of SOCE by NO and others claim the contrary. In addition, several different mechanisms of action are proposed.

In one of the first reports, Bischof et al. reported that the NO donor sodium nitroprusside (SNP) at 30 µM induced inhibition of SOCE while lower 0.5 µM SNP had an opposite effect and led to modest enhancement of the Ca²⁺ entry [203]. Similar findings about the dual, concentration dependent effects of NO and NOdonors on SOCE were also reported in neutrophils and platelets in 2011 [204,205]. Bolotina and co-workers showed in two successive reports that NO inhibits SOCE in human platelets and proposed an accelerated SERCA pumping rate and consequently faster refilling of the ER-Ca²⁺ stores as the molecular mechanism underlying this event [206,207]. That NO suppresses SOCE function in HEK 293 cells was also shown by Groschner and his colleagues. This effect was induced by application of either NO donor (2-(N,Ndiethylamino)diazenolate-2-oxide sodium salt, DEANO) or pure NO $(100 \,\mu\text{M})$. Interestingly, the authors could show that SOCE sensitivity towards NO is abolished upon overexpression of TRPC3 channels [208]. Several additional reports also showed inhibition of SOCE by NO, proposing different mechanisms [209-212].

It has also been reported that NO can have stimulatory effects on SOCE [213]. In 1998 and 1999, the group of Gill reported that NO donors GEA3162, SNP and NO_2 activated SOCE independently of guanylate cyclase and suggested thiol oxidation as a modulating mechanism. According to their data, the authors suggest S-nitrosylation either on the proteins within the ER stores or on the PM channels as an underlying mechanism of NO induced SOCE activation [214,215]. These findings were confirmed by Watson et al. who proposed that NO induces depletion of ryanodine-sensitive Ca^{2+} stores and thus leads to activation of SOCE [216].

In 2003, Moneer et al. showed differential NO sensitivity of two independent Ca²⁺ entry pathways activated by vasopressin in a vascular smooth muscle cell-line. Generation of NO following vasopressin treatment was shown to activate a non-capacitative Ca²⁺ entry pathway, while inhibiting a capacitative Ca²⁺ entry via the protein kinase G [217]. Inhibition of SOCE by an NO donor was also reported by Watson at al. [218]. A summary of redox regulation of CRAC/Orai channels is given in Table 4.

6. Implications and future perspectives

Regulation of proteins by reversible or irreversible oxidation, glutathionylation or nitrosylation is becoming an increasingly recognized aspect of cellular signaling. The currently used approaches often involve external addition of oxidants. Given the variable and often contradictory results it is questionable to what extent our experimental designs can mimic conditions of ROS and RNS production and diffusion in a biological system. The short-lived nature of these reactive species makes accurate quantification of physiologically produced concentrations a very difficult task.

Many proteins contain a number of potentially reactive cysteines, hindering the interpretation of the effects of ROS in the absence of mutagenesis studies. While mutagenesis of specific residues often is able to pinpoint the molecular targets of ROS (e.g. C157 in TRPV1), this is not always the case, possibly when cysteines

can be in a reactive and in a non-reactive state (i.e. C558 in TRPC5) or when mutation of a particular cysteine leads to non-functional channels. Future work is also needed to clearly separate effects of potentially increased phosphorylation due to inactivation of phosphatases. The development of faster and more accurate in vivo ROS indicators will increase elucidation of physiological ROS production and its effects. Considering the important roles of Ca²⁺ conducting ion channels such as CaV, TRP and CRAC/Orai channels for cellular functions, detailed molecular analyses of their redox regulation on a molecular and physiological level will help to understand the interplay between Ca²⁺ and ROS signaling. The many inconsistent findings regarding ROS modulation of Ca²⁺ channels highlight the necessity to better control and quantify ROS, to identify molecular targets and to consider complex interactions between them.

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