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# Transit of H<sub>2</sub>O<sub>2</sub> across the endoplasmic reticulum membrane is

## not sluggish

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#### Abstract

Cellular metabolism provides various sources of hydrogen peroxide ( $H_2O_2$ ) in different organelles and compartments. The suitability of  $H_2O_2$  as an intracellular signaling molecule therefore also depends on its ability to pass cellular membranes. The propensity of the membranous boundary of the endoplasmic reticulum (ER) to let pass  $H_2O_2$  has been discussed controversially. In this essay, we challenge the recent proposal that the ER membrane constitutes a simple barrier for  $H_2O_2$  diffusion and support earlier data showing that (i) ample  $H_2O_2$  permeability of the ER membrane is a prerequisite for signal transduction, (ii) aquaporin channels are crucially involved in the facilitation of  $H_2O_2$  permeation, and (iii) a proper experimental framework not prone to artifacts is necessary to further unravel the role of  $H_2O_2$ permeation in signal transduction and organelle biology.

#### Principles of reduction-oxidation signaling

The life of a multicellular organism is organized in a complex network of intercellular communication. In this vein, individual cells react to external cues such as hormones or other receptor-based agonists by the activation of signal transduction cascades, which faithfully transfer the extracellular signals to the intracellular addressees. Similar processes are activated also in single cell organisms in response to pheromones or nutrient signals. The single steps of

these signaling cascades are designed to proceed by optimized spatial and temporal dynamics [1].

An important element in intracellular signal transduction is the transient formation of diffusible second messengers, which allow amplification of the signal due to their multiple places of action. Amongst many other second messengers, hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) is now widely being recognized to serve as such a mobile signaling molecule [2, 3]. H<sub>2</sub>O<sub>2</sub> is one of the reactive oxygen species that are produced upon reduction of molecular oxygen and is itself an oxidant. It primarily acts by specifically oxidizing target proteins on specialized, sensitive cysteine residues to modulate their function [4]. Therefore, H<sub>2</sub>O<sub>2</sub>-mediated signaling is referred to as reduction-oxidation (redox) signaling. Of note, H<sub>2</sub>O<sub>2</sub> is a relatively poorly reactive oxidant, which allows it to travel further from its site of generation than can superoxide (O<sub>2</sub><sup>...</sup> ) or hydroxyl radical, before it encounters a peroxidase, catalase or signaling target [4, 5].

A prime example of redox signaling is the role of  $H_2O_2$  during growth factor-stimulated signal transduction [6, 7]. Here, the binding of extracellular growth factor ligands to receptor tyrosine kinases (RTKs) on the cell surface frequently co-activates members of the NADPH oxidase (Nox) family [8-10]. Nox family members locally produce  $O_2^{\cdots}$ , which rapidly dismutates to  $H_2O_2$ . This increase in  $O_2^{\cdots}$  and  $H_2O_2$  generation is required for sustained receptor tyrosine phosphorylation and downstream signaling events, because  $H_2O_2$  inactivates protein tyrosine phosphatases (PTPs) on a reactive cysteine in their active site [11, 12]. Membrane topology is a critical aspect in this process. The  $O_2^{\cdots}$  -producing active sites of Nox complexes are located on the exoplasmic side of the membrane, whereas PTPs localize to the cytosol. This topological problem is solved by membrane-embedded aquaporin channels (AQPs). By serving as  $H_2O_2$  pores they facilitate the formation of local areas with an elevated  $H_2O_2$  concentration on both sides of the plasma membrane [13-16].

#### $H_2O_2$ can readily permeate through the endoplasmic reticulum membrane

RTK signaling is not restricted to the plasma membrane. For instance, epidermal growth factor (EGF) receptor can be internalized upon stimulation by endocytosis and brought into proximity with the endoplasmic reticulum (ER) membrane [17]. As a notable consequence, ER-associated proteins such as Nox4 [18, 19] and the phosphatase PTP1b [20-22] play important roles during EGF receptor signaling by acting in analogy to their cognate signaling components at the plasma membrane [17]. Nox4, which can directly, i.e. irrespective of a dismutase, generate  $H_2O_2$  in the ER lumen [23-27], is coupled to the transient oxidative inactivation of PTP1b on the cytosolic side of the ER membrane [17]. This sequence of events premises that  $H_2O_2$  must be able to pass the ER membrane at a time scale that copes with EGF receptor signaling.

While observations of redox signaling at the ER are relatively scarce at this stage, it is clear that  $H_2O_2$  is widely utilized as a signaling molecule *in vivo* [28] and it is quite predictable that further mechanisms specific to the ER will be uncovered in the future [18]. Other examples, which are connected to  $H_2O_2$  transit across the ER membrane, are granulocyte colony-stimulating factor receptor signaling [29], oxidative DNA damage in response to cellular stresses [30-32], activation of survival pathways upon  $H_2O_2$  generation in the ER [33, 34], and the regulatory roles of ER-luminal peroxidases in various settings of cytosolic signal transduction [29, 35-38]. These findings clearly indicate the permeability of the ER membrane for  $H_2O_2$ .

Ample  $H_2O_2$  permeability at the ER membrane has additionally been demonstrated by studying over-expressed ER oxidoreductin 1 $\alpha$  (Ero1 $\alpha$ ). This ER-luminal oxidase produces  $H_2O_2$ , which is immediately detoxified by the Ero1 $\alpha$ -associated peroxidase GPx8 [18]. Depletion of GPx8, however, leads to the overflow of  $H_2O_2$  to the cytosol [39]. By contrast,

depletion of the ER-luminal high-abundance-high-affinity-high-turnover-peroxidase peroxiredoxin 4 [40, 41] does not cause similar leakage of Ero1 $\alpha$ -derived H<sub>2</sub>O<sub>2</sub> into the cytosol [39]. Thus, the shielding of the cytosol against Ero1 $\alpha$ -derived H<sub>2</sub>O<sub>2</sub> takes place at the Ero1 $\alpha$ -GPx8 interface through catalytic elimination [42]. If hindered diffusion of H<sub>2</sub>O<sub>2</sub> at the ER membrane was to provide an additional shielding mechanism, Ero1 $\alpha$ -derived H<sub>2</sub>O<sub>2</sub> would certainly be eliminated by peroxiredoxin 4 already within the ER and not found in the cytosol upon depletion of GPx8.

### Aquaporins regulate the permeability of the ER membrane to $H_2O_2$

Is the transport of  $H_2O_2$  at the ER facilitated by AQPs in analogy to the situation at the plasma membrane? AQP8 fulfills a major function in the transport of  $H_2O_2$  at the plasma membrane [13]. In addition, knockdown of AQP8 strongly diminishes the entry of exogenous  $H_2O_2$  into the ER of plasma membrane-permeabilized cells [13]. This indicates that AQP8 can accelerate the transit of  $H_2O_2$  also across the ER membrane when expressed at physiological levels. Since cell surface AQP8 is synthesized at the ER before trafficking to the plasma membrane, a physiological function in the ER is conceivable. This is also supported by its steady-state localization both at the plasma membrane and in "intracellular vesicles" [43]. In addition, AQP8 appears to be involved in the transit of  $H_2O_2$  from mitochondria in certain cell types [44].

AQP8 and other AQPs show specific tissue distributions. The rich collection of human AQPs enables a versatile regulation of transmembrane permeation of water throughout the body by harboring specific differences in transcriptional regulation, post-translational modification, protein stability, water permeability, and subcellular distribution [43]. Accordingly, it is likely that AQPs other than AQP8 play complementary, tissue- and context-specific roles with

regard to  $H_2O_2$  transport at the ER. One obvious candidate is AQP11, the subcellular localization of which is strongly shifted to the ER [45, 46]. AQP11 loss-of-function causes destructive symptoms of ER stress, which mainly manifest in the proximal tubular epithelial cells of the kidney [45-47] but also in other organs such as the liver [48]. The failure of AQP11-deficient cells is accompanied by elevated levels of intracellular H<sub>2</sub>O<sub>2</sub> [45]. Whether or not ER stress and H<sub>2</sub>O<sub>2</sub> dysregulation are linked to a change in H<sub>2</sub>O<sub>2</sub> permeability of the ER membrane remains to be shown.

In addition to the tissue-specific expression level of ER AQPs, the  $H_2O_2$  permeability of the ER membrane is likely regulated by post-translational modifications. For instance, the permeability of AQP8 is reversibly inhibited in response to diverse stress conditions through the targeting of cysteine 53 (Iria Medraño-Fernandez, Stefano Bestetti, and R.S.; unpublished observations) and the overproduction of ER-luminal  $H_2O_2$  appears to stimulate its own passage through the ER membrane in liver cells of living mice [49].

Based on biophysical and structural data, it has been deduced that all AQPs that are able to transport water can also transport  $H_2O_2$  [50]. Thus, not only the highly conducting aquaammoniaporin AQP8 but also the water-permeable AQP11 is predicted to serve as a *bona fide*  $H_2O_2$  channel.

#### The ER membrane is not refractory to rapid $H_2O_2$ diffusion

In a recent publication, the ER membrane was postulated to comprise a significant barrier to  $H_2O_2$  diffusion [51]. This postulate was based on an experiment, in which oxidation of intracellular  $H_2O_2$  probes in response to increasing concentrations of extracellular  $H_2O_2$  were recorded. As already worked out elsewhere [39], the  $H_2O_2$ -dependent oxidation of the genetically encoded probe HyPer [52] was recorded upon concomitant addition of the

disulfide reductant dithiothreitol (DTT). In this setup, ER-targeted HyPer was less readily oxidized than cytosolic HyPer [51]. This appears to be a trivial observation though, as exogenous  $H_2O_2$  on its way to the ER must cross the cytosol, which is equipped with a plethora of powerful peroxidases. In a comparable experimental setup, most  $H_2O_2$  was consumed before it could reach the depth of the cell [53]. Konno et al. addressed this issue by using a cell line that expresses relatively low levels of some cellular antioxidant enzymes [51], a measure that can modulate but not eliminate the problem of cytosolic dissipation of  $H_2O_2$ . The less efficient oxidation of ER-targeted HyPer compared to cytosolic HyPer therefore cannot only be interpreted to reflect hampered permeability of the ER membrane to  $H_2O_2$ .

In addition to cytosolic and ER-targeted HyPer, Konno et al. used mitochondrial HyPer, which showed similar H<sub>2</sub>O<sub>2</sub>-induced fluorescence changes as cytosolic HyPer [51]. This is surprising, because, as for the ER, mitochondria can only be reached via the cytosol, which would be expected to decrease the H<sub>2</sub>O<sub>2</sub>-sensitivity of mitochondrial HyPer below the sensitivity of cytosolic HyPer (see above). How can this be explained? HyPer is not only sensitive to oxidation but also to alkalinisation [52], which is typically controlled for by also analyzing the response of cysteine-mutant HyPer [54]. Of potential relevance, treatment of cells with  $H_2O_2$  induces the transient alkalinisation of the mitochondrial matrix [55]. Furthermore, we note that the responses to extracellular H<sub>2</sub>O<sub>2</sub> of chemical, pH-independent H<sub>2</sub>O<sub>2</sub> sensors are similarly slow in mitochondria and ER and slightly faster in cytosol and nucleus [56]. Apart from pH, other organelle-specific differences in the handling of HyPer could also be relevant. It is possible, for example, that the rich collection of thiol-disulfide isomerases in the ER (for review see [23, 57]) catalyzes the reduction of ER-targeted HyPer by DTT particularly well. This in turn would decrease the net steady-state oxidation of ERtargeted HyPer as compared to mitochondrial HyPer at the lower doses of H<sub>2</sub>O<sub>2</sub>, as has been observed [51]. Although these explanations are yet hypothetical, we suggest that some

mitochondrion-specific feature rather than the relative impermeability of the ER membrane causes the more pronounced response to  $H_2O_2$  of mitochondrial HyPer compared to ER-targeted HyPer.

In summary, all published data strongly support the notion that facilitated permeability to

H<sub>2</sub>O<sub>2</sub> is a designated and likely regulated feature of the ER membrane, which is in line with

the central signaling role of this fascinating organelle.

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#### Highlights

- Ample H<sub>2</sub>O<sub>2</sub> permeability of the ER membrane is critical for signal transduction
- Aquaporins facilitate the transmembrane permeation of H<sub>2</sub>O<sub>2</sub>
- The ER H<sub>2</sub>O<sub>2</sub> pool appears not to be isolated from other cell compartments