

1 Peroxisomes contribute to intracellular calcium dynamics

2 Yelena Sargsyan¹, Uta Bickmeyer¹, Katrin Streckfuss-Bömeke^{2,4}, Ivan Bogeski³, Sven Thoms^{1,4*}

3 ¹Department of Child and Adolescent Health, University Medical Center, Göttingen, Germany

4 ²Clinic for Cardiology and Pneumology, University Medical Center, Göttingen, Germany

5 ³Molecular Physiology, Institute of Cardiovascular Physiology, University Medical Center, Göttingen,
6 Germany

7 ⁴German Center of Cardiovascular Research (DZHK), Partner site Göttingen, Germany

8 * Correspondence: sven.thoms@med.uni-goettingen.de

9 **Keywords:** peroxisomes, calcium, genetically encoded calcium indicator (GECI), cardiomyocytes, iPSC

10 Abstract

11 Peroxisomes communicate with other cellular compartments by transfer of various metabolites. However,
12 whether peroxisomes are sites for calcium handling and exchange has remained contentious. Here we
13 generated sensors for assessment of peroxisomal calcium and applied them for single cell-based calcium
14 imaging in HeLa cells and cardiomyocytes. We found that peroxisomes in HeLa cells take up calcium upon
15 depletion of intracellular calcium stores and upon calcium influx across the plasma membrane. Further, we
16 show that peroxisomes of neonatal rat cardiomyocytes and human induced pluripotent stem cell-derived
17 cardiomyocytes can take up calcium in a controlled manner. Our results indicate that peroxisomal and
18 cytosolic calcium signals are tightly interconnected. Hence, peroxisomes may play an important role in
19 shaping cellular calcium dynamics by serving as buffers or sources of intracellular calcium.

20 Introduction

21 Calcium ions (Ca^{2+}) play a decisive role in the regulation of many cellular processes and inter-compartment
22 communication, especially in excitable cells like neurons or cardiomyocytes (CMs) (Clapham, 2007). In CMs,
23 for example, cytosolic Ca^{2+} directly engages in cell contraction. At the same time, mitochondrial Ca^{2+}
24 coordinates ATP production and energy demand in CMs (Williams et al., 2015), highlighting the importance
25 of intracellular organelles in Ca^{2+} redistribution. The main sites of Ca^{2+} entry to the cell and intracellular
26 calcium signal regulation are the plasma membrane (PM) and intracellular calcium stores, in particular those
27 of the endoplasmic reticulum (ER) (Paupe and Prudent, 2018).

28 Excess of organellar Ca^{2+} can be detrimental for health. Elevated mitochondrial uptake increases
29 mitochondrial reactive oxygen species (ROS) production and is associated with heart failure and ischemic
30 brain injury (Santulli et al., 2015; Starkov et al., 2004). Reversely, mitochondrial ROS decreases if Ca^{2+} uptake
31 to mitochondria is suppressed (Mallilankaraman et al., 2012; Tomar et al., 2016). Understanding of principles
32 and mechanisms of organellar Ca^{2+} handling provides starting point to develop interventions in dysregulated
33 calcium handling.

34 Peroxisomes are small intracellular organelles with a phospholipid bilayer membrane. In concert with
35 evolutionarily conserved functions in lipid and ROS metabolism, peroxisomes are highly plastic and change in
36 their number, morphology and content upon environmental stimuli (Smith and Aitchison, 2013).
37 Communication of peroxisomes with other cellular compartments through exchange of ROS or lipid
38 metabolites is essential for human health (Castro et al., 2018; Schrader et al., 2020; Wanders et al., 2015).
39 Yet, peroxisomal Ca^{2+} has not been studied in excitable cells before, and there are contradicting data about
40 the Ca^{2+} handling in peroxisomes and its dependence on cytosolic Ca^{2+} (Drago et al., 2008; Lasorsa et al.,
41 2008). It has been suggested that peroxisomes are potential targets of Ca^{2+} signaling pathways that initiate
42 outside of the peroxisome or serve as a cytosolic Ca^{2+} buffer, but peroxisomes may also take up Ca^{2+} due to
43 their own need (Drago et al., 2008; Islinger et al., 2012).

44 Measurement of Ca^{2+} dynamics *in vivo* inside cellular organelles was driven by the development of Ca^{2+} -
45 sensitive fluorescent proteins, also known as genetically encoded Ca^{2+} indicators (GECIs) (Gibhardt et al.,
46 2016; Pozzan and Rudolf, 2009). Ca^{2+} dynamics was analysed in the ER, in mitochondria, the cytosol, and in
47 lysosomes by using GECIs (McCue et al., 2013; Whitaker, 2010). GECIs have a Ca^{2+} binding domain, usually
48 calmodulin (CaM). Ratiometric pericam is a single fluorophore-based GECI with circularly permuted EYFP
49 (cpEYFP) as the fluorophore (Nagai et al., 2001). A special role among GECIs play chameleon-based sensors
50 that use Förster resonance energy transfer (FRET). Here, Ca^{2+} results in a conformational change, that
51 decreases the distance between donor (typically CFP) and acceptor (typically a YFP variant) so that FRET
52 occurs (Gibhardt et al., 2016; Palmer and Tsien, 2006; Pérez Koldenkova and Nagai, 2013). The FRET/donor
53 ratio (FRET ratio) correlates with the Ca^{2+} concentration (Figure 1A).

54 The possibility to generate human induced pluripotent stem cells (hiPSCs) from somatic cell sources and to
55 direct their differentiation into almost any cell type make it possible to maintain and study human CMs in
56 culture (Yoshida and Yamanaka, 2011).

57 This work combines the advantages of organelle-targeted GECIs and hiPSCs. We develop several peroxisomal
58 Ca^{2+} sensors, and we measure intraperoxisomal Ca^{2+} after pharmacological stimulation in non-excitable and
59 excitable cells. We show that peroxisomes take up Ca^{2+} upon cytosolic Ca^{2+} increase both following ER Ca^{2+}
60 store depletion and Ca^{2+} entry to the cells through PM. We also demonstrate that peroxisomes take up Ca^{2+}
61 in rat CMs and hiPSC-CMs.

62 Results

63 Development and validation of Ca^{2+} sensors for peroxisomal Ca^{2+}

64 To assess peroxisomal Ca^{2+} , we used three GECIs with different affinities to Ca^{2+} : D3cpV, D1cpV and pericam
65 (Figure 1A, Table 1). The sensors were chosen to cover a wide range of K_d values to identify the most suitable
66 GECI for intra-peroxisomal measurement. We preferred ratiometric sensors that allow measurements in two
67 wavelengths. This enables direct interpretation of the acquired data by calculating the ratio of intensities at
68 each time point. The ratios provide direct information about Ca^{2+} binding and are independent of the sensor
69 concentration itself (Pérez Koldenkova and Nagai, 2013). For the direct comparison of cellular
70 compartments, we used specific sensors for the cytosol (D3cpV, R-GECO1), mitochondria (4mtD3cpV), and
71 peroxisomes (D3cpV-px, D1cpV-px, pericam-px) (Figure 1B).

72 D3cpV is a chameleon-type indicator based on FRET. The conformational change associated with the Ca^{2+}
73 binding to CaM leads to an increase in FRET efficiency and FRET ratio (Pérez Koldenkova and Nagai, 2013).

74 D3cpV has a K_d value of 0.6 μM and a dynamic range of 5.0 (Palmer et al., 2006). D1cpV, in comparison, is a
75 FRET sensor with a K_d value of 60 μM (Palmer et al., 2004). Finally, pericam is a cpEYFP-based GECl with two
76 excitation peaks at ~ 420 nm and 505 nm (Nagai et al., 2001). In the presence of Ca^{2+} , a conformational
77 change in the pericam structure shifts the excitation profile so that the 505/420 ratio increases and serves as
78 a measure of Ca^{2+} concentration (Figure 1A). Pericam has a K_d value of 1.7 μM and dynamic range of 10.

79 We added strong peroxisomal targeting signals of the PTS1 type to D3cpV, D1cpV, and pericam and tested
80 their localization after transfection by co-staining with antibodies directed against the peroxisomal
81 membrane protein PEX14. All constructs targeted to peroxisomes (Figure 1C).

82 To test in living cells if D3cpV-px senses Ca^{2+} in the peroxisome, we permeabilized cells by digitonin, washed
83 out the cytosol, and added relatively high Ca^{2+} concentrations. Ca^{2+} addition resulted in drastic increase of
84 FRET and a 1.5-fold increase in FRET ratio (Figure 1D). In order to further illustrate the increase of the FRET
85 signal, we false-colored by using a color look-up table (LUT) the images recorded before and after Ca^{2+} the
86 addition (Figure 1D).

87 When we performed the same type of experiment with D1cpV-px, FRET increased as well after Ca^{2+} addition,
88 showing that the D1cpV-px construct is Ca^{2+} sensitive (Figure 1E). However, following the same stimulation
89 protocol, the signal change of D1cpV-px was only 1.08-fold, and thus considerably smaller than with using
90 D3cpV-px. Due to the low signal change we excluded D1cpV-px from the further experiments on peroxisomal
91 Ca^{2+} . Using pericam-px, the third peroxisome-targeted sensor in our experiments, high concentration of Ca^{2+}
92 addition after digitonin treatment resulted in 1.5-fold increase similar as for D3cpv-px (Figure 1F). Based on
93 these results we decided to use D3cpv-px and pericam-px to evaluate Ca^{2+} dynamics in peroxisomes in
94 further experiments.

95 To study possible mislocalisation or residual signal of peroxisomal Ca^{2+} sensors from the cytosol, we again
96 analysed peroxisomal Ca^{2+} signals following digitonin stimulation of intact cells. In the case of mislocalisation
97 of the sensor to cytosol the signal decrease after digitonin stimulation is expected. We first tested this in
98 D3cpV-px (Figure 1G). There was no signal change observed, suggesting that D3cpV-px has no cytosolic
99 mislocalisation. The cytosol washout also did not change the Ca^{2+} signal of the pericam-px before and after
100 digitonin wash, suggesting that pericam-px, like D3cpV-px, is exclusively localized to the peroxisome (Figure
101 1H).

102 **Measurement of peroxisomal Ca^{2+} in non-excitabile cells**

103 We first aimed to compare the maximal possible response of cytosol and peroxisomes to Ca^{2+} . On that
104 purpose, we used ionomycin as an ionophore. Ionomycin resulted in fast and immediate increase of cytosolic
105 signal (Figure 2A). Peroxisomal signal increased also, yet, gradually. After reaching its maximum it decreased
106 gradually and in 12 minutes almost returned to its starting values. The cytosolic reached its half maximal
107 value in the same time with most significant decrease observed in the first two minutes after the maximum.
108 This observations suggest that there could also be differences in Ca^{2+} handling also under near-physiological
109 stimulation.

110 Based on the Ca^{2+} measurements in other organelles (Matsuda et al., 2013; Petrunaro et al., 2015; Suzuki et
111 al., 2014; Zhao et al., 2011) we developed an experimental paradigm for peroxisome responses to the
112 depletion and refilling of intracellular Ca^{2+} stores, specifically ER, in non-excitabile HeLa cells (Figure 2B). The
113 stimulation of cell-surface localized G-protein coupled receptors by 100 μM histamine results in the
114 activation of phospholipase C cascade. Inositol 1,4,5-trisphosphate (IP_3), the product of the cascade, binds to
115 the IP_3 receptor on the ER membrane, triggering Ca^{2+} store release. The cells are then exposed to 1 mM

116 extracellular Ca^{2+} , which leads to store-operated Ca^{2+} entry (SOCE) and a second Ca^{2+} elevation in the cytosol.
117 Ca^{2+} is constantly pumped back to the ER through sarcoplasmic/endoplasmic reticulum calcium ATPase
118 (SERCA) (Clapham, 2007).

119 When we treated HeLa cells expressing D3cpV-px according to this protocol, we observed two peaks (Figure
120 2C). Histamine addition resulted in a steep and fast increase of intraperoxisomal Ca^{2+} based on depletion of
121 the ER. Addition of extracellular Ca^{2+} resulted in more gradual increase and gradual return to basal levels
122 (Figure 2C).

123 Using the measurements with D3cpV-px and the known properties of the sensor, we calculated the absolute
124 Ca^{2+} concentration (Figure 2D) applying the formula described by Palmer and Tsien (2006). Under basal
125 conditions, Ca^{2+} level in peroxisomes is around 400 nM and it rises upon near-physiological stimulation with
126 histamine up to 1.8 μM Ca^{2+} (Figure 2E). The Ca^{2+} dynamics in peroxisomes measured with D3cpV-px was
127 reproduced by pericam-px: a larger peak is observed after ER-store depletion and a smaller one after
128 extracellular Ca^{2+} addition. The observed ratio curve from pericam-px largely resembles that from D3cpV-px.
129 Since pericam has a K_d value of 1.7 μM and covers higher Ca^{2+} concentrations, the observed result confirms
130 the upper limit of peroxisomal Ca^{2+} and the range of Ca^{2+} between 0.4 and 1.8 μM (Figure 2F). However,
131 pericam is described as pH sensitive (Nagai et al., 2001), and since there is currently no consensus regarding
132 pH levels in peroxisomes (Dansen et al., 2000; Jankowski et al., 2001; Waterham et al., 1990) we decided to
133 perform all further experiments with D3cpV-px.

134 To confirm that the response in our experiments is due to the immediate increase in Ca^{2+} concentration, and
135 to be able to directly compare peroxisomal Ca^{2+} handling with that of the cytosol, cells were co-transfected
136 with D3cpV-px and the mApple-based cytosolic Ca^{2+} sensor R-GECO1, that increases in intensity when
137 binding Ca^{2+} (Zhao et al., 2011). A large increase in the red signal from R-GECO1 was observed both upon ER
138 store depletion and addition of extracellular Ca^{2+} (blue curve in Figure 2G). Although the GECIs used for the
139 measurement in two compartments have different properties that can result in differences in their kinetics,
140 peroxisomes largely follow the Ca^{2+} changes in the cytosol. Interestingly, there is little or no delay between
141 signal increase in cytosol and peroxisomes, and the post-stimulation decline is more gradual and prolonged
142 in peroxisomes compared to the cytosol, indicating the existence of a possible barrier or gate that can be
143 saturated (Figure 2H).

144 To compare peroxisomal Ca^{2+} levels at rest and under stimulation with that of cytosol and mitochondria, cells
145 were transfected with D3cpV sensors targeting specifically these compartments. FRET ratio was assessed as a
146 direct indicator of Ca^{2+} concentration (Figure 2I-L). All three compartments showed two peaks: one after ER-
147 store depletion with histamine, and another after extracellular Ca^{2+} addition and PM-based uptake (Figure
148 2I).

149 The basal levels of calcium in mitochondria and peroxisomes detected with this sensor were comparable and
150 significantly higher than that in the cytosol (typically ≈ 100 nM, Paupe and Prudent, 2018) in the current
151 settings (Figure 2J). Furthermore, the increase of Ca^{2+} in peroxisomes upon intracellular store depletion with
152 100 μM histamine was significantly lower than the increase in the cytosol or mitochondria (Figure 2K),
153 speaking against the hypothesis that peroxisomal Ca^{2+} is rising drastically upon stimulation as suggested
154 before (Lasorsa et al., 2008). The addition of extracellular Ca^{2+} resulted in another peak in all three
155 compartments (Figure 2L), evidencing that peroxisomes, like mitochondria depend on the PM-based uptake.
156 Altogether, this suggests that peroxisomes tend to follow Ca^{2+} dynamics of the cytosol.

157 **Peroxisomal Ca²⁺ measurement in cardiomyocytes**

158 We decided to test in neonatal rat cardiomyocytes (NRCMs) the hypothesis that Ca²⁺ can access cardiac
159 peroxisomes. NRCMs are primary cells with a well-developed T-tubule system and serve as a model for
160 electrophysiological studies on CMs (Soeller and Cannell, 1999; Morad and Zhang, 2017).

161 We adapted the chemical stimulation protocol for the CMs by reducing it to a single stimulation, since the
162 main source of Ca²⁺ in these cells is the ER. We used thapsigargin (Tg) to chemically stimulate the CMs (Figure
163 3A). Tg is a SERCA antagonist and blocks the constant repumping of Ca²⁺ back to the ER, resulting in Ca²⁺
164 accumulation in the cytosol. To avoid measurement distortion due to the spontaneous contractile activity of
165 CMs, they were treated with 2,3-butanedione monoxime (BDM) (Gwathmey et al., 1991) before the
166 experiment. As a proof of concept and for direct comparison, we performed the first round of measurements
167 using the cytosol-localized Ca²⁺-sensor D3cpV (Figure 3B-D). A comparison between the Tg-treated cells with
168 the buffer conditions (Figure 3B) demonstrated, as expected, no differences in the basal ratios (Figure 3C),
169 but an increase of cytosolic Ca²⁺ upon Tg addition (Figure 3D).

170 To measure peroxisomal Ca²⁺ changes, we transfected NRCMs with D3cpV-px and compared Tg treatment
171 with the untreated control group (Figure 3E). No offset of basal ratios between the two groups was present
172 before treatment (Figure 3F). After the addition of the SERCA inhibitor, peroxisomal Ca²⁺ increased,
173 evidencing peroxisomal Ca²⁺ uptake in NRCMs after store depletion (Figure 3G).

174 In the next set of experiments, we wanted to know if peroxisomes of human cardiac cells are able to take up
175 Ca²⁺. To test this, human iPSCs created from fibroblasts of a healthy donor were differentiated into CMs
176 using standardized protocols including cardiac mesoderm induction by subsequent activation and inhibition
177 of the WNT pathway (Lian et al., 2013) and metabolic selection (Tohyama et al., 2013) (Figure 3H). Cardiac
178 differentiation was tested for homogeneity by using the cardiac specific marker cardiac troponin T (cTNT)
179 and analysis by flow cytometry at day 90 of differentiation. Our differentiation consisted of 90 %-95 % cTNT-
180 positive cells (Figure 3I). Staining of hiPSC-CMs with antibodies against α -actinin showed a regular sarcomeric
181 striation pattern (Figure 3J).

182 As a proof of concept and for direct comparison, we measured cytosolic and peroxisomal Ca²⁺ and compared
183 Tg treatment with the addition of Ca²⁺-free buffer without Tg to the control cells (Figure 3K-P). Starting with
184 the same basal ratios as the control samples (Figure 3K and 3L), Tg-treated cells showed a Ca²⁺ increase after
185 the treatment (Figure 3M).

186 After confirming that Tg can effectively deplete Ca²⁺ stores in hiPSC-CMs, we measured peroxisomal Ca²⁺ in
187 these cells (Figure 3N). No ratio differences were present before Tg treatment (Figure 3O). Ca²⁺-store
188 depletion resulted in an increase of peroxisomal Ca²⁺, confirming peroxisomal Ca²⁺ uptake in hiPSC-CMs
189 (Figure 3P).

190 To test whether Ca²⁺ enters peroxisomes in a beat-to-beat manner in NRCMs, we field stimulated the cells
191 with 1 Hz frequency (Figure 4A-D). The action potential depolarizes cell membrane resulting in the activation
192 of voltage-gated Ca²⁺ channels in T-tubules (Bootman et al., 2002; Chapman, 1979). As a result, initial minor
193 amount of Ca²⁺ enters the cell. It activates ryanodine receptors on the sarcoplasmic reticulum membrane,
194 resulting into Ca²⁺ release from the stores. Ca²⁺-induced Ca²⁺ release from the stores enables cardiac muscle
195 contraction. During relaxation SERCA and NCX (sodium-calcium exchanger) pump Ca²⁺ back to the
196 intracellular Ca²⁺ stores and out of the cells (Clapham, 2007).

197 Under field stimulation, we observed rhythmical changes of Ca^{2+} level in the cytosol (Figure 4A). To quantify
198 the amplitude of changes and link to the stimulation we performed fast Fourier transformation (FFT) of the
199 data (Figure 4B). Signal amplitude oscillations in the cytosol were rhythmical and corresponded to the
200 stimulation frequency (Figure 4B).

201 To test peroxisomal response to electrical stimulation, NRCMs expressing D3cpV-px were paced at a
202 frequency of 1 Hz (Figure 4C). Oscillations observed were smaller in amplitude and appeared less regular
203 than the cytosolic responses. To identify the frequency domain of these oscillations we performed FFT
204 (Figure 4D). The extracted pattern showed amplitude changes at 1 Hz, suggesting that peroxisomes take up
205 Ca^{2+} in beat-to-beat manner.

206 Altogether, these results suggest that peroxisomes in both, rat and human cardiomyocytes were able to take
207 up Ca^{2+} upon intracellular Ca^{2+} -store depletion and cytosolic Ca^{2+} increase.

208 Discussion

209 Peroxisomes are metabolically highly active organelles in need of communication with other cellular
210 compartments (Sargsyan and Thoms, 2020). ROS signaling and homeostasis are central to the participation of
211 peroxisomes in signaling pathways (Lismont et al., 2019). In the current work we focused on Ca^{2+} dynamics of
212 peroxisomes as one of the major signaling molecules in the cell. We demonstrate that Ca^{2+} can enter
213 peroxisomes of HeLa cells both when ER-stores are depleted and when cytosolic Ca^{2+} increases after Ca^{2+}
214 entry across the plasma membrane.

215 Two articles published in 2008 brought forth conflicting data on peroxisomal Ca^{2+} . According to Drago et al.
216 (2008), the basal level of Ca^{2+} in peroxisomes equals the cytosolic Ca^{2+} level, whereas Lasorsa et al. (2008)
217 find peroxisomal Ca^{2+} to be 20 times higher than in the cytosol. While Lasorsa et al. (2008) report rise of
218 peroxisomal Ca^{2+} up to 100 μM using an aequorin-based sensor, Drago et al. (2008) suggest slow increase
219 when cytosolic Ca^{2+} rises. Each of the groups used a single yet different technique. These differences in the
220 results can be partially attributed to the different measurement methods and the cell types used. Aequorin
221 imaging requires long incubation times and cell population-based analysis that can be disadvantageous when
222 measuring Ca^{2+} in intracellular organelles. In our experiments with HeLa cells, we found four-fold higher basal
223 peroxisomal Ca^{2+} level compared to the cytosol and increase up to 1.8 μM upon stimulation (Table 1). The
224 range of the changes we report are based on the measurements with D3cpV-px and are supported by the
225 measurement with pericam-px. Hence, we conclude that D3cpV-px can be used for measuring peroxisomal
226 Ca^{2+} concentration in a broad variety of cell types.

227 Electron microscopic experiments on rodent hearts performed in the 1970s show that peroxisomes are
228 closely associated with T-tubules and with junctional sarcoplasmic reticulum (Hicks and Fahimi, 1977). The
229 sarcoplasmic reticulum is an indispensable site for the excitation-contraction coupling and Ca^{2+} handling in
230 myocytes (Flucher et al., 1994). The localization of peroxisomes to these sites raises the question if cardiac
231 peroxisomes react to Ca^{2+} oscillations on a beat-to-beat basis, and/or if they can buffer calcium. HiPSC-CMs
232 provide a wide spectrum of possibilities in cardiac research ranging from drug screening to cardiac
233 regeneration (Yoshida and Yamanaka, 2011). In addition, these cells have been especially used to study
234 patient-specific disease models including arrhythmic disorders and cardiomyopathies demonstrating a robust
235 correlation to the predicted phenotype (Borchert et al., 2017; Liang et al., 2016; Streckfuss-Bömeke et al.,
236 2017). We report here that Ca^{2+} is entering peroxisomes upon intracellular Ca^{2+} -store depletion in CMs. Since

237 intracellular store depletion is the main source of Ca^{2+} in CMs in the process of excitation-contraction
238 coupling, it can be hypothesized that peroxisomes take up Ca^{2+} also on beat-to-beat manner in these cells.

239 Measurement of peroxisomal Ca^{2+} in CMs with FRET sensors in field stimulation suggests that peroxisomal
240 Ca^{2+} increases on beat-to-beat manner. This suggests that peroxisomes may participate in excitation-
241 contraction processes. The exact role of peroxisomes here is the matter of future research. Furthermore, the
242 experimental protocols developed here can be applied to study peroxisomal Ca^{2+} other cell types like
243 neurons.

244 We found that basal peroxisomal Ca^{2+} levels are higher than cytosolic levels. There are two major ways of
245 generating this Ca^{2+} gradient on the two sides of the membrane. One option could be the energy-dependant
246 uptake mechanism, like SERCA for the ER (Clapham, 2007). We are, however, not aware of any data that can
247 support this model. The second option may be locally high Ca^{2+} concentration at the entry side that would
248 allow more direct channeling of Ca^{2+} (from the ER) into the peroxisomes resulting in relatively high
249 peroxisomal Ca^{2+} . This second mechanism is known from the mitochondrial Ca^{2+} handling, where ER-
250 mitochondria contact sites with tethering proteins generate microdomain with locally high Ca^{2+}
251 concentration (Hirabayashi et al., 2017). As a result, Ca^{2+} entry to mitochondria follows the Ca^{2+} gradient but
252 mitochondrial Ca^{2+} is higher compared to cytosol. For the plausibility of the second option for peroxisomes
253 speak the existance of ER-peroxisome contact sites (Costello et al., 2017; Hua et al., 2017). Therefore, we
254 propose a hypothetical model of this mechanism, where most of the components are, however, yet
255 unknown (Figure 5).

256 Although peroxisomal Ca^{2+} levels are higher than cytosolic Ca^{2+} levels, peroxisomes are unlikely to store
257 significant amounts of Ca^{2+} under normal conditions, and they themselves take up Ca^{2+} when intracellular
258 stores are depleted. Under specific conditions, like apoptosis or oxidative stress, the situation may change,
259 however. We show that the rise of peroxisomal Ca^{2+} after histamine stimulation is not delayed and largely
260 follows the cytosolic Ca^{2+} . Though there could be a delay due to the binding and conformational changes of
261 GECs needed before the detection of the increase of the FRET signal, the range of this delay is less than
262 milliseconds and cannot be seen in the experiments described here. We conclude that peroxisomes respond
263 to cytosolic Ca^{2+} since we only found concordant changes of Ca^{2+} concentration in these two compartments.

264 The question of the cellular function and potential targets of peroxisomal Ca^{2+} is still open. One of the roles of
265 Ca^{2+} could be the regulation of peroxisomal processes. On the other hand, metabolic processes themselves
266 may regulate Ca^{2+} uptake to organelles, as known from mitochondria (Nemani et al., 2020). Whether there is
267 a mutual regulation of metabolic pathways or ROS production localised to peroxisomes is not known. Some
268 plant but not mammalian catalases bind Ca^{2+} (Yang and Poovaiah, 2002). Currently, there are no peroxisomal
269 processes known in mammals that would depend on Ca^{2+} . Peroxisomes, however, could serve as an
270 additional cytosolic buffer for Ca^{2+} to take up an excess of cytosolic Ca^{2+} and release it slowly. Based on the
271 findings of this study that the Ca^{2+} concentration in the peroxisome is higher than in the cytosol, it could be
272 that peroxisomes may also serve as additional Ca^{2+} source for the cytosol in extreme situations.

273 **Methods**

274 **DNA constructs**

275 D3cpV-px (PST 1738) was generated from (pcDNA-)D3cpV (kind gift from A. Palmer and R. Tsien (Palmer et
276 al., 2006) (Addgene #36323)) by amplifying an insert with OST 1599 (GCGCATCGAT GGTGATGGCC
277 AAGTAACTA TGAAGAG) and OST 1600 (GCGCGAATTC TTAGAGCTTC GATTTCAGAC TTCCTCGA) primers. The
278 product was then reinserted into D3cpV using Clal and EcoRI restriction sites. (pcDNA-)4mtD3cpV was a kind
279 gift from A. Palmer and R. Tsien (Palmer et al., 2006) (Addgene #36324). D1cpV-px (PST 2169) was generated
280 from the (pcDNA-) D1cpV (Palmer et al., 2004) (Addgene #37479) by amplifying an insert with
281 oligonucleotide OST 2003 (GCGCGATCC CATGGTGAGC AAGGGC) and OST 2002 (CGCGGAATTC TTAGAGCTTC
282 GATTTCAGAC TTCCTATGAC AGGCTCGATG TTGTGGCGGA TCTGAAGTT). The product was then reinserted
283 into D1cpV using EcoRI and BamHI restriction sites. Pericam-px (PST 2170) was generated from ratiometric-
284 pericam (for mitochondria) (Nagai et al., 2001) by amplifying an insert with OST 2116 (GCGCAAGCTT
285 ATGAAGAGGCGC TGAAGAAAA) and OST 2117b (GCGCGAATTC CTAGAGCTTC GATTTCAGAC TTCCTATGAC
286 AGGCTTTGCT GTCATCATTT GTACAACT), which was then re-inserted into ratiometric-pericam using EcoRI
287 and HindIII restriction sites. (CMV-)R-GECO1 was a kind gift from R. Campbell (Zhao et al., 2011).

288 **Cells, cell culture and immunofluorescence**

289 HeLa cells were cultured in low glucose Dulbecco's Modified Eagle Medium (DMEM) medium (Biochrom)
290 supplemented with 1% Pen/Strep (100units/ml Penicillin and 100 μ g/ml Streptomycin), 1% (w/v) glutamine
291 and 10% (v/v) Fetal Calf Serum (FCS) in 5% CO₂ at 37 \square C. For immunofluorescent detection of PEX14, cells
292 were fixed with 4% paraformaldehyde for 20 min, and permeabilized using 0.5% Triton X-100 in PBS for 5
293 min. After blocking for 30 min with 10% BSA in PBS (blocking buffer) at 37 \square C, antigens were labelled with
294 primary antibodies at 37 \square C for 1 h. Rabbit anti-PEX14 (ProteinTech) primary antibody dilution in blocking
295 buffer was 1:500. Labeling with the secondary antibodies conjugated to Cy3 (Life Technologies) was done for
296 1 h (1:500). Coverslips were mounted with ProLong Gold mounting medium with or without DAPI (Thermo
297 Fisher Scientific).

298 NRCMs were isolated from newborn rats. Briefly, after the rats were sacrificed hearts were removed from
299 the thoracic cavity, homogenized mechanically and digested in 1mg/ml collagenase type II containing calcium
300 and magnesium-free PBS at 37 \square C with magnetic stirring. Supernatant was taken every 20 min and transferred
301 to DMEM medium supplemented with Glutamax (Thermo Fisher Scientific), FCS and 1% Pen/Strep. Cells
302 were then centrifuged, the cell pellet resuspended in fresh medium and transferred to a Petri dish for 45 min
303 (37 \square C and 5% CO₂). The fibroblasts adhered and NRCMs remained in the supernatant. NRCMs were then
304 seeded on glass cover slips covered by Geltrex (Thermo Fisher Scientific).

305 Cells and cardiac differentiation of hiPSCs were described earlier (Borchert et al., 2017). Cells were studied 90
306 days after initiation of differentiation. Following differentiation, purity of hiPSC-CMs was determined by flow
307 analysis (>90% cardiac TNT⁺) or by morphology (Borchert et al., 2017). HiPSC-CMs were maintained in RPMI
308 1640 supplemented with Glutamax, HEPES and B27 supplement.

309 **Ca²⁺ measurements**

310 Cells (200,000 for HeLa and hiPSC-CMs and 500,000 for NRCMs) were seeded on glass cover slips and
311 transfected with sensor plasmids using Effectene (Qiagen) (HeLa) or Lipofectamine LTX Reagent (Thermo
312 Fisher Scientific) (hiPSC-CMs and NRCMs) according to the manufacturer's instructions. Cells were imaged
313 using a Zeiss Observer D1 (equipped with Zeiss Colibri 2 and Evolve 512 Delta EMCCD acquisition camera) or
314 Axio Observer Z1 (equipped Zeiss Colibri 7, Definite Focus.2 and Zeiss AxioCam 702) with 40 \times oil Fluar (N.A.
315 1.3) objective at 37 \square C in a Ca²⁺-free imaging buffer (145 mM NaCl, 4 mM KCl, 10 mM HEPES, 10 mM glucose,

316 2 mM MgCl₂, 1 mM EGTA, pH 7.4 at 37°C) 24 hours (HeLa and NRCMs) or 48 hours (hiPSC-CMs) after
317 transfection. Where indicated, NRCMs were field-stimulated at 1 Hz with MyoPacer ES (IonOptix). Data were
318 analyzed with AxioVision (Zeiss) and ZEN (Zeiss) software. Background and bleed-through (BT) were
319 corrected in the FRET/donor ratio:

$$\frac{FRET}{donor} = \frac{(FRET - background) - [(CFP - background) \times BT] - [(YFP - background) \times BT]}{CFP - background}$$

320 Excitation 420 nm and 505 nm with emission filters 483 ± 16 nm and 542 ± 14 nm, or excitation 438 ± 12 nm
321 and 508 ± 11 nm with emission filters 479 ± 20 nm and 544 ± 14 nm were used. Where indicated, the
322 concentration of Ca²⁺ in the imaging buffer was increased to 1 mM by doubling the buffer volume to the cells
323 (e.g. during treatment with chemicals) by the addition of Ca²⁺-containing buffer (imaging buffer that contains
324 2 mM CaCl₂ (pH 7.4, 37°C) instead of EGTA). HiPSC-CMs and NRCMs were incubated in 10 mM 2,3-
325 butanedione monoxime (BDM) before the measurements. FRET ratios (calculated as FRET donor ratio) were
326 calculated by subtracting the background intensity and correcting for crosstalk. ER-store depletion in the
327 cells was induced by 100 μM histamine (HeLa) in Ca²⁺-free buffer or 1 μM Tg in Ca²⁺-containing buffer. For
328 permeabilization, cells were treated with 0.01% digitonin in Ca²⁺-free EGTA buffer for 50 sec to 1 min and
329 cytosol was washed out by rinsing twice with Ca²⁺-free EGTA buffer. Cell response to ionomycin was
330 measured by the addition of 5 μM ionomycin in 10 mM Ca²⁺-containing buffer. Images for color LUT were
331 made by applying Royal LUT on difference image of FRET and CFP in case of D3cpV-px and D1cpV-px, or
332 difference image of 505 nm and 420 nm in case of pericam-px.

333 Statistical analysis

334 Statistical significance was assessed using two-sided unpaired student's t-test when comparing two groups,
335 or one-way ANOVA followed by Tukey's post hoc test when three groups were compared. Data were
336 presented as Tukey's box plots: the box is limited by 25th and 75th percentiles. Data points larger than 75th
337 percentile plus 1.5IQR (interquartile range) or smaller than 25th percentile minus 1.5IQR are presented as
338 outliers. All other data are covered by the whiskers.

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448 **Author Contributions**

449 ST conceived and designed the study. YS performed most experiments, conducted data analysis and
450 prepared all figures. UB tested D3cpV-px localization, and measured Ca^{2+} in NRCMs and hiPSC-CMs. IB
451 supervised Ca^{2+} measurements and provided access to the Zeiss Cell Observer Z1 with Colibri3 LED system.
452 KSB provided hiPSC-CMs and contributed to manuscript writing. YS and ST wrote the manuscript. All authors
453 read, revised and approved the manuscript.

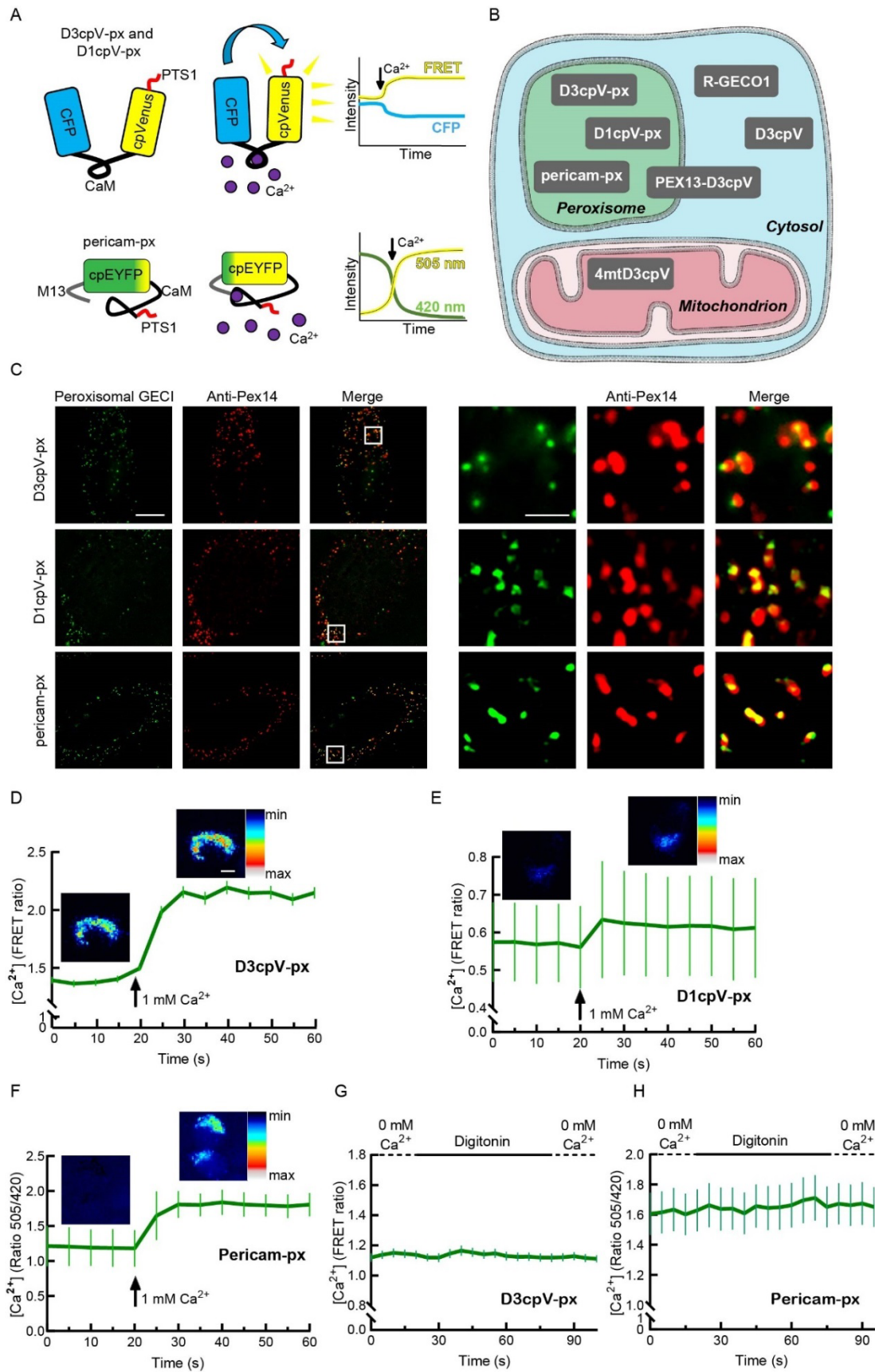
454 **Funding**

455 This project was supported by grants from the Deutsche Forschungsgemeinschaft TH 1538/3-1 to ST, the
456 Collaborate Research Council 'Modulatory units in heart failure' SFB 1002/2 TP A10 to ST and SFB1190 TP17
457 and SFB1027 TP C4 to IB, the MWK/VW foundation Project 131260 /ZN2921 to ST, the Horst and Eva-Luise
458 Köhler Foundation to ST, the Fritz Thyssen Foundation Az 10.19.2.026MN to KSB, and a PhD stipend by the
459 DAAD program 57381412 ID 91572398 to YS.

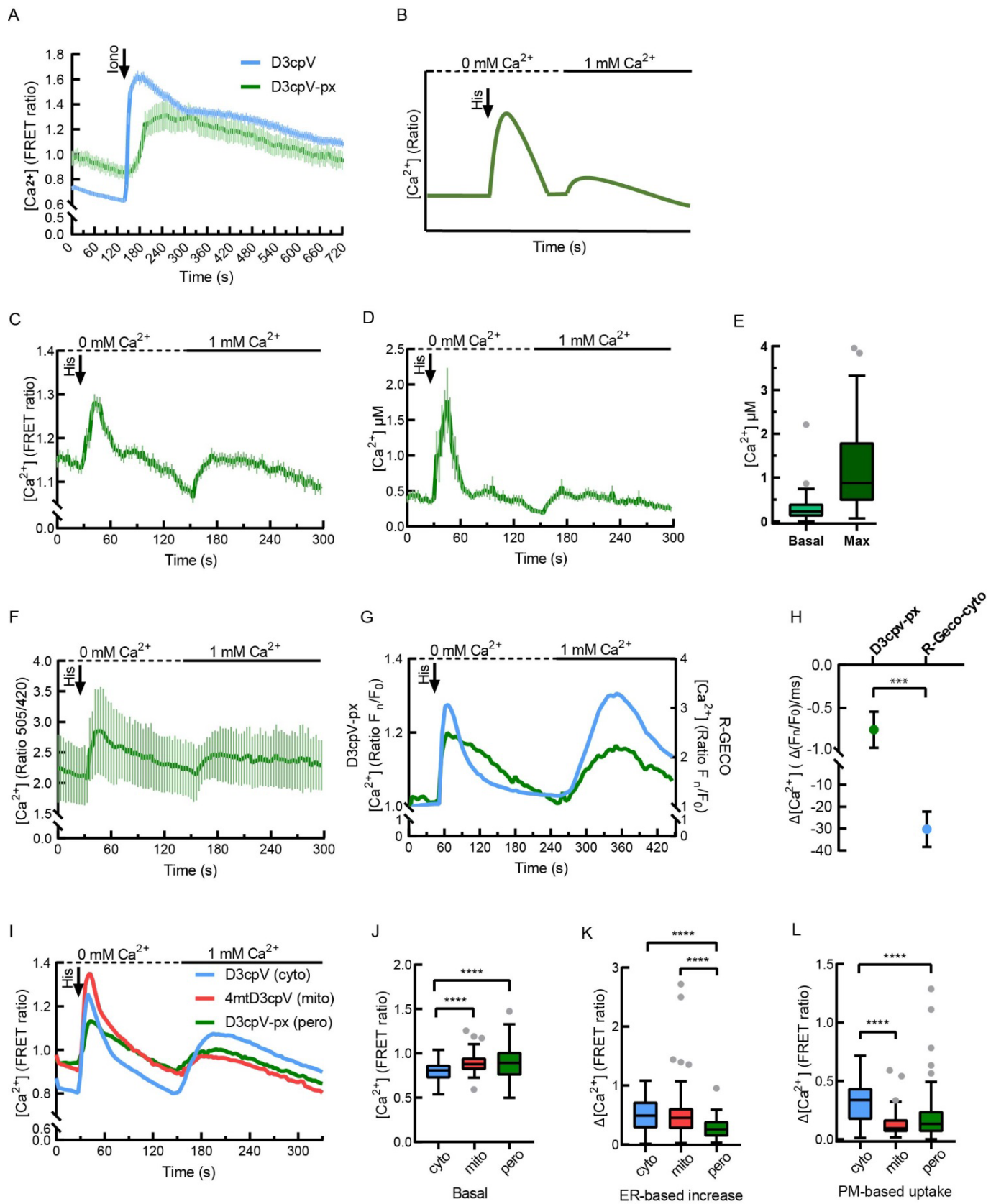
460 **Acknowledgments**

461 We thank Drs. Robert Campbell, Takeharu Nagai, and Nicolas Demaurex for providing GECO and pericam
462 plasmids. We thank Julia Hofhuis for earlier work on peroxisomal calcium and for cloning of D3cpV-px and
463 Xin Zhang for support with microscopy.

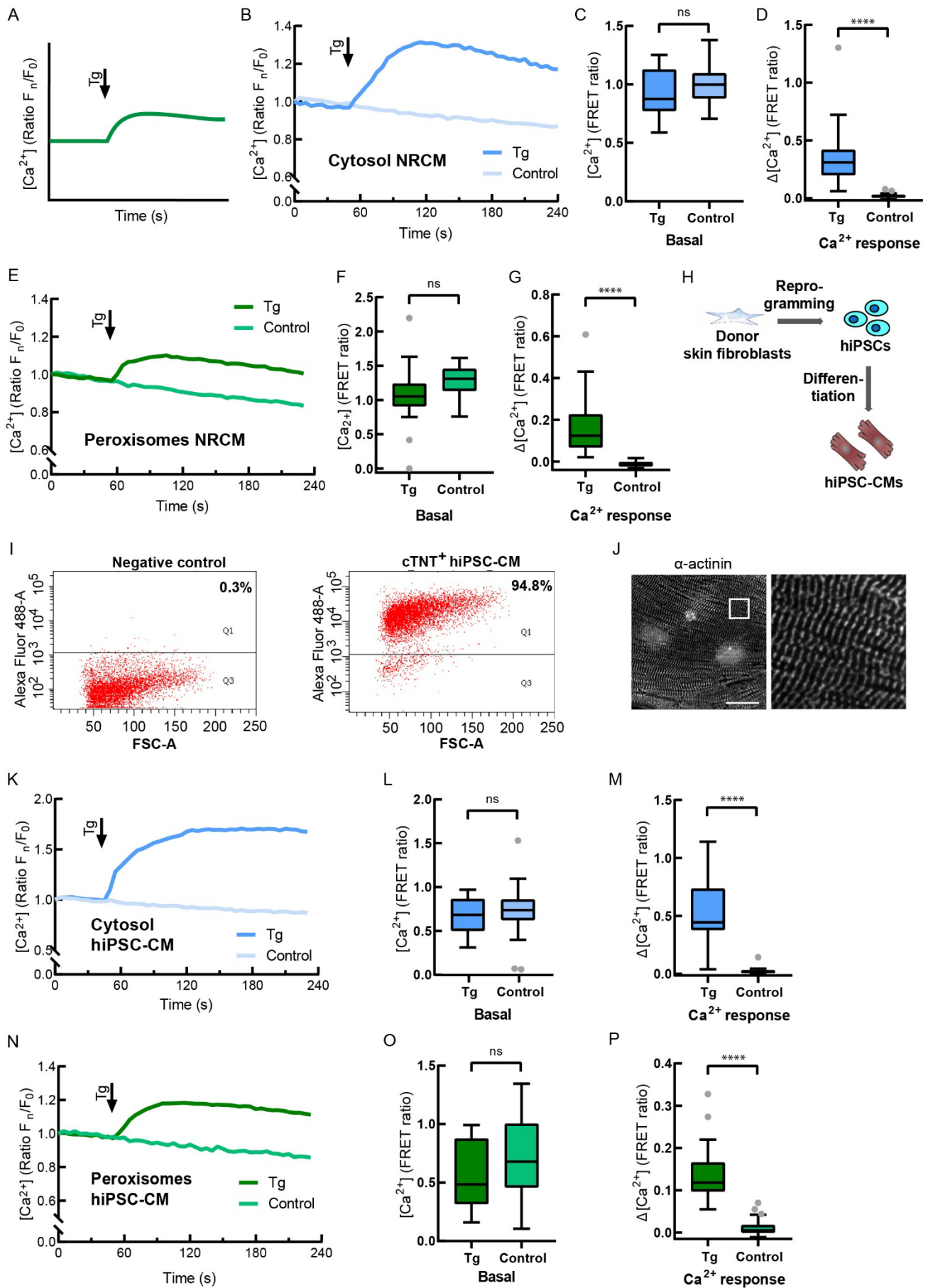
464 **FIGURES**



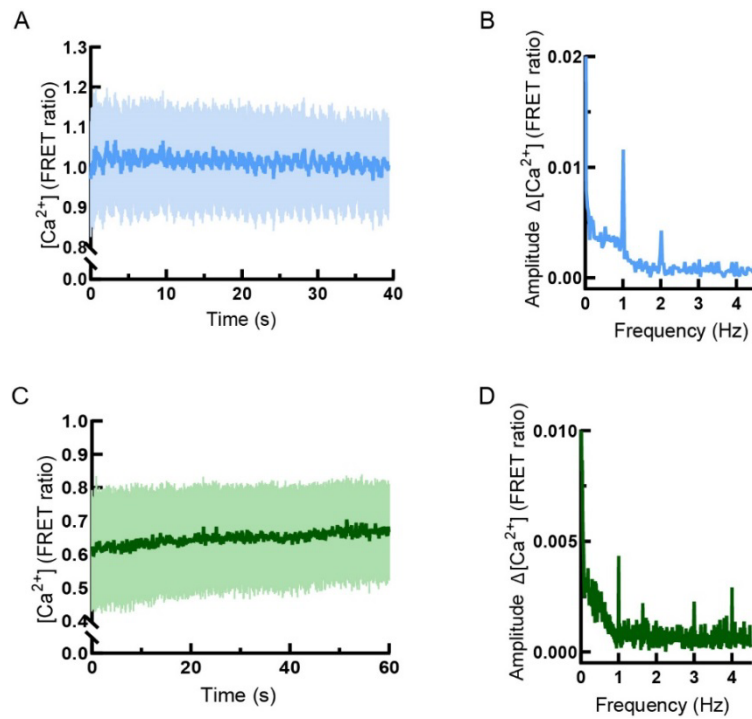
466 **Figure 1: New sensors for peroxisomal Ca²⁺.** (A) Genetically encoded calcium indicators (GECIs) targeted to
467 peroxisomes. D3cpv-px and D1cpV-px are FRET sensors with modified CaM sites. Pericam-px is a single-
468 fluorophore based GECI that has M13 and CaM as Ca²⁺ binding sites. In the absence of Ca²⁺, the emission
469 measured when the sensor is excited with 420 nm is higher than when excited with 505 nm. The ratio
470 505/420 is a measure for the Ca²⁺ concentration. (B) Subcellular localization of GECIs used in this study. (C)
471 Peroxisomal GECIs colocalize with the peroxisomal membrane marker PEX14. HeLa cells were transfected
472 with the GECIs and stained with anti-PEX14 antibodies. The images in the left part of the panel show one cell
473 each (scale bar 10 μm). The cropped areas are marked and magnified in the right part of the panel (scale bar
474 2 μm). (D-F) D3cpv-px, D1cpv-px, and pericam-px are Ca²⁺ sensitive. Images false-colored with LUT show
475 representative cells before (left) and after (right) Ca²⁺ addition. Curves presented as mean ± SEM. Scale bar:
476 10 μm. (D) Addition of 1 mM Ca²⁺ to D3cpV-px expressing cells results in 1.5-fold FRET ratio increase, n = 60
477 cells from three independent experiments. (E) FRET ratio increases 1.08 times when 1 mM Ca²⁺ is added to
478 D1cpv-px expressing cells, n = 33 cells from three experiments. (F) Ca²⁺ addition leads to 1.5-fold increase in
479 505/420 ratio with Pericam-px, n = 75 cells from three experiments. (G) Measurement of D3cpV-px during
480 cytosol washout. No change in signal is detected. (H) Measurement of pericam-px during cytosol washout.
481 No difference of signal before and after cytosol washout is detected, n = 43 cells for D3cpV-px in (G) and n =
482 45 cells for pericam-px in (H).



484 **Figure 2: Measurement of peroxisomal Ca^{2+} in HeLa cells.** (A) Comparison of cytosolic and peroxisomal
485 responses to ionomycin (Iono). In comparison to cytosol, peroxisomal signal increases gradually, $n = 16$ cells
486 for D3cpV and $n = 9$ cells for D3cpV-px. (B) Experimental paradigm of a two-step Ca^{2+} measurement in non-
487 excitable cells. 1st peak after histamine (His) addition: ER-store depletion. 2nd peak, after addition of
488 extracellular Ca^{2+} : PM-based uptake. (C) Measurement with D3cpV-px according to the paradigm in (B). Two
489 Ca^{2+} peaks of the experimental paradigm are detectable with D3cpV-px, $n \geq 50$ cells from three experiments.
490 (D) Absolute Ca^{2+} concentration dynamics calculated from the data in (C). (E) Basal and maximum (max) Ca^{2+}
491 concentrations in peroxisomes based on (C). (F) Measurement with pericam-px according to the paradigm in
492 (B), $n = 27$ cells from three experiments. (G) Simultaneous measurement of cytosolic (blue) and peroxisomal
493 (green) Ca^{2+} . No delay of signal increase after histamine addition, but a delayed drop of the signal in
494 peroxisomes. Left y axis: of D3cpV-px (peroxisomal sensor). Right y axis: F_n/F_0 ratio of R-GECO1 (cytosolic
495 sensor), $n = 35$ cells from three experiments. (H) Decline of F_n/F_0 ratio per millisecond (ms) in the linear part
496 of the curves in (G) (from second 65 to 115, Student's t-test). Kinetic delay in decrease in peroxisomal signal
497 is seen. (I) Comparison of cytosolic, mitochondrial and peroxisomal Ca^{2+} response measured following the
498 paradigm in (B). Characteristic two peaks present in all three compartments. (J) Basal levels of Ca^{2+} in
499 peroxisomes are similar to mitochondria. Analysis performed based on the data from (I). (K) Peroxisomal Ca^{2+}
500 increase upon ER-store depletion is smaller than that of cytosol or mitochondria. Analysis performed based
501 on the data from (I). (L) Peroxisomal Ca^{2+} increase upon PM-based cellular uptake of Ca^{2+} is comparable to
502 mitochondria. Analysis performed based on the data from (I). (A, C, D, F) Data presented as mean \pm SEM. (J-
503 L) One-way ANOVA followed by Tukey's post hoc test was used for the statistical analysis. *** $p < 0.001$,
504 **** $p < 0.0001$, Tukey's box plots. Cyto: cytosolic, mito: mitochondrial, pero: peroxisomal. $n = 83$ (cyto), 116
505 (mito), 117 (pero) cells from six independent experiments.

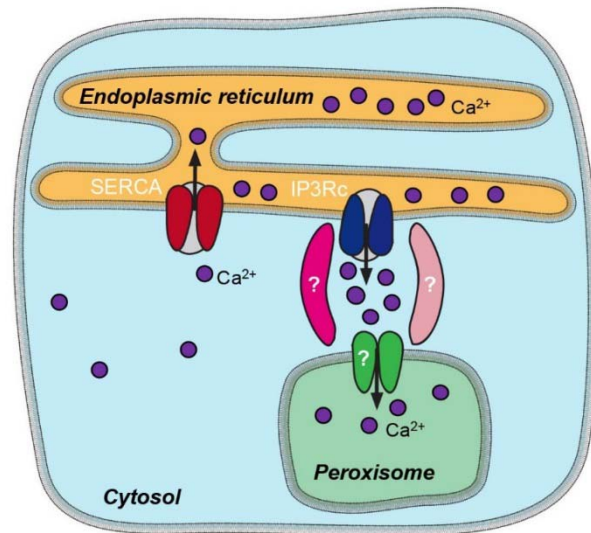


507 **Figure 3: Measurement of peroxisomal Ca^{2+} in cardiomyocytes.** (A) Experimental paradigm of Ca^{2+}
508 measurement in excitable cells. The peak after thapsigargin (Tg) addition represents Ca^{2+} increase due to the
509 SERCA inhibition and Ca^{2+} retention in the cytosol. (B) Cytosolic Ca^{2+} measurement in NRCMs following the
510 experimental design in (A), n = 25 (Tg), 22 (control) from three experiments. Addition of Tg is compared to
511 the addition of Tg-free buffer (control). (C) Basal levels are not different before the treatment in (B). (D) After
512 Tg addition in (B) cytosolic Ca^{2+} increases. (E) Peroxisomal Ca^{2+} measurement in NRCMs following the
513 experimental design in (A). Addition of Tg is compared to the addition of Tg-free buffer (control), n = 20 (Tg),
514 31 (control) from three experiments. (F) Basal levels of Ca^{2+} are not different before the treatment in (E). (G)
515 Peroxisomal Ca^{2+} increases after Tg addition in (E). (H) hiPSC-CMs generation. Donor skin fibroblasts were
516 reprogrammed to hiPSCs, which were then differentiated to CMs. (I) hiPSC-CMs were stained for cardiac
517 troponin (cTnT) and analyzed by flow cytometry. Negative control without primary antibody. 94.8% of iPSC-
518 CMs are cTnT-positive (cTnT⁺). (J) Immunofluorescence staining visualized α -actinin protein expression and
519 regular sarcomeric organization. Scale bar: 20 μm . (K) Cytosolic Ca^{2+} measurement in hiPSC-CMs with D3cpV
520 following the experimental paradigm for excitable cells in (A). Addition of Tg is compared to the addition of
521 Tg-free buffer (control) to avoid artefacts and false results due to mechanical effect on the cells due to the
522 addition itself. n = 24 (Tg), 27 (control) from three differentiation experiments. (L) No difference is found
523 between two groups before the treatment in (K). (M) Tg addition in (K) results in cytosolic Ca^{2+} increase. (N)
524 Peroxisomal Ca^{2+} measurement in hiPSC-CMs with D3cpV-px following the experimental design in for
525 excitable cells depicted in (A). Addition of Tg is compared to the addition of Tg-free buffer (control). n = 26
526 (Tg), 33 (control) from three differentiation experiments. (O) Basal levels of Ca^{2+} are not different before the
527 treatment in (N). (P) Peroxisomal Ca^{2+} increases after Tg addition in (M). (B, E, K, N) Data presented as means
528 from three independent experiments. (C, D, F, G, L, M, O, P) Unpaired Student's t-test was used for the
529 statistical analysis. ****p < 0.0001, Tukey's box plots.



530

531 **Figure 4: Measurement of peroxisomal Ca^{2+} in paced cardiomyocytes.** (A) D3cpV transfected NRCMs are
532 stimulated with 1 Hz. Images are taken every 50 ms. Oscillations of FRET ratio are seen, $n = 3$. (B) FFT from
533 the data in (A). Signal increases are rhythmic and correspond to the pacing frequency. (C) D3cpV-px
534 transfected NRCMs are stimulated with 1 Hz. Images are taken every 100 ms. FRET ratio oscillations are seen.
535 $n = 3$. (D) FFT from the data in (C). Signal increases are rhythmic and correspond to the pacing frequency.



536

537 **Figure 5: Peroxisomal Ca²⁺ entry and cellular Ca²⁺ distribution.** ER Ca²⁺ release triggers Ca²⁺ entry into the
 538 peroxisome. In our hypothetical model, ER-peroxisome proximity defines Ca²⁺ microdomains with locally
 539 elevated Ca²⁺ concentration shielded from the cytosol. As a result, Ca²⁺ entry to peroxisomes follows the local
 540 gradient but peroxisomal Ca²⁺ is eventually higher than in the cytosol. IP3Rc: IP₃ receptor calcium release
 541 channel of the ER.

542 **Table**

543 **Table 1. Key properties of the GECIs for cytosol and peroxisome**

Cytosolic GECIs			Peroxisomal GECIs (this study)	
Construct Name	K _d (<i>in vitro</i>)	Dynamic range, D	Construct Name	Maximal increase upon 1 mM Ca ²⁺ addition
D3cpV	0.6 μM ⁽¹⁾	5.0 ⁽¹⁾	D3cpV-px	1.50 x
D1cpV	60 μM ⁽²⁾	1.7 ⁽³⁾	D1cpV-px	1.08 x
Ratiometric-pericam	1.7 μM ⁽⁴⁾	10.0 ⁽⁴⁾	Pericam-px	1.50 x

544 References: ¹ Palmer et al. (2006); ² Palmer et al. (2004); ³ Greotti et al. (2016) ⁴ Nagai et al. (2001)