



Calibration and functional analysis of three genetically encoded Cl⁻/pH sensors

M. Mukhtarov^{1,2}, L. Liguori³, T. Waseem^{1,4}, F. Rocca^{3,5}, S. Buldakova¹, D. Arosio^{3*} and P. Bregestovski^{1*}

¹ Brain Dynamics Institute, Inserm UMR1106, Aix-Marseille University, Marseille, France

² Laboratory of Neurobiology, Institute of Fundamental Medicine and Biology, Kazan Federal University, Kazan, Russia

³ Institute of Biophysics, National Research Council and FBK, Trento, Italy

⁴ Institute of Biophysics and Cell Engineering, National Academy of Sciences, Minsk, Belarus

⁵ Physics Department, University of Trento, Trento, Italy

Edited by:

Pau Gorostiza, Catalan Institution for Research and Advanced Studies, Spain

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Colin J. Akerman, University of Oxford Academic Research Fellow, UK

Alexandre Mourot, Université Pierre et Marie Curie, France

*Correspondence:

D. Arosio, Institute of Biophysics, National Research Council and FBK, Trento, Italy.

e-mail: daniele.arosio@cnr.it;

P. Bregestovski, Brain Dynamics Institute, Inserm UMR1106, Aix-Marseille University, 27 Boulevard Jean Moulin, 13385 Marseille Cedex 05, France.
e-mail: piotr.bregestovski@univmed.fr

Monitoring of the intracellular concentrations of Cl⁻ and H⁺ requires sensitive probes that allow reliable quantitative measurements without perturbation of cell functioning. For these purposes the most promising are genetically encoded fluorescent biosensors, which have become powerful tools for non-invasive intracellular monitoring of ions, molecules, and enzymatic activity. A ratiometric CFP/YFP-based construct with a relatively good sensitivity to Cl⁻ has been developed (Markova et al., 2008; Waseem et al., 2010). Recently, a combined Cl⁻/pH sensor (ClopHensor) opened the way for simultaneous ratiometric measurement of these two ions (Arosio et al., 2010). ClopHensor was obtained by fusion of a red-fluorescent protein (DsRed-monomer) to the E²GFP variant that contains a specific Cl⁻-binding site. This construct possesses pK_a = 6.8 for H⁺ and K_d in the 40–50 mM range for Cl⁻ at physiological pH (~7.3). As in the majority of cell types the intracellular Cl⁻ concentration ([Cl⁻]_i) is about 10 mM, the development of sensors with higher sensitivity is highly desirable. Here, we report the intracellular calibration and functional characterization of ClopHensor and its two derivatives: the membrane targeting PalmPalm-ClopHensor and the H148G/V224L mutant with improved Cl⁻ affinity, reduced pH dependence, and pK_a shifted to more alkaline values. For functional analysis, constructs were expressed in CHO cells and [Cl⁻]_i was changed by using pipettes with different Cl⁻ concentrations during whole-cell recordings. K_d values for Cl⁻ measured at 33°C and pH ~7.3 were, respectively, 39, 47, and 21 mM for ClopHensor, PalmPalm-ClopHensor, and the H148G/V224L mutant. PalmPalm-ClopHensor resolved responses to activation of Cl⁻-selective glycine receptor (GlyR) channels better than did ClopHensor. Our observations indicate that these different ClopHensor constructs are promising tools for non-invasive measurement of [Cl⁻]_i in various living cells.

Keywords: fluorescent biosensors, intracellular chloride, intracellular pH, non-invasive monitoring, optogenetics

INTRODUCTION

Beginning with the pioneer studies by Tsien and co-authors on measurements of intracellular calcium concentration in intact lymphocytes (Tsien et al., 1982), non-invasive monitoring of different intracellular ions (Ca²⁺, Mg²⁺, Cl⁻, and H⁺) has become a powerful direction of research for functional analysis of neurons and other cell types under normal and pathophysiological conditions.

Among several methods proposed for monitoring intracellular Cl⁻ concentration ([Cl⁻]_i), the most promising is the use of genetically encoded Cl⁻-sensitive probes (Bregestovski et al., 2009; Mancuso et al., 2011). The first generation of probes named “Clomeleon” (Kuner and Augustine, 2000) and “Cl-Sensor” (Markova et al., 2008) was based on the Cl⁻-sensitive yellow-fluorescent protein (YFP) linked with Cl⁻-insensitive

cyan-fluorescent protein (CFP), which was the reference fluorescence molecule. These indicators have opened the way for non-invasive monitoring and ratiometric measurement of [Cl⁻]_i in different cell types *in vitro* (Pellegrino et al., 2011; Bertolini et al., 2012; Friedel et al., 2013). Expressed in neurons of transgenic mice, they have allowed imaging of Cl⁻ dynamics in inhibitory circuits of different brain areas, including hippocampus, cerebellum, and deep cerebellar nuclei (Berglund et al., 2008, 2011), as well as in intact hippocampus (Dzhala et al., 2012) and a dorsal root ganglia preparation (Batti et al., 2013). Producing a construct consisting of a glycine receptor (GlyR) with Cl⁻-Sensor incorporated into the long cytoplasmic domain (BioSensor-GlyR) provided a tool for non-invasive monitoring activity of these Cl⁻-selective receptor-operated channels (Mukhtarov et al., 2008).

The recently proposed combined Cl⁻/pH sensor (ClopHensor) opened the way for simultaneous ratiometric measurement of these two ions (Arosio et al., 2010). ClopHensor is obtained by fusion of a red-fluorescent protein (DsRed-monomer) to the E²GFP variant that contains a specific Cl⁻-binding site. This construct is particularly promising as it allows simultaneous monitoring of [Cl⁻]_i and intracellular pH (pH_i) in the same cells (Arosio et al., 2010; Bregestovski and Arosio, 2011; Raimondo et al., 2012). At physiological pH (~7.3), ClopHensor possesses pK_a = 6.8 for H⁺ and K_d in the 40–50 mM range for Cl⁻. As in the majority of cell types the intracellular Cl⁻ concentration is about 10 mM (Khirug et al., 2008; Tyzio et al., 2008; Bregestovski et al., 2009), development of sensors with higher sensitivity is necessary.

Here, we present calibration and functional analysis of ClopHensor and two of its derivatives: (i) PalmPalm-ClopHensor, which should have preferable membrane targeting and thus allow near-membrane measurement of [Cl⁻]_i and pH_i; and (ii) ClopHensor with mutations H148G and V224L aimed to increase the affinity of this probe for Cl⁻ and change the pH-sensing properties. It has been previously demonstrated that the V224L mutation increases the affinity for Cl⁻ by an order of magnitude (Arosio et al., 2011), while the H148G mutation increases the pK_a value by 1 pH unit in YFP (Elslinger et al., 1999) as well as in GFP (Hanson et al., 2002). Here, we combined the H148G and the V224L mutations in E²GFP in order to: (1) shift the K_d for Cl to the lower mM range, and (2) increase the pK_a of the GFP sensing element to more alkaline values.

For these three constructs, calibrations determining their sensitivity to Cl⁻ and H⁺ in living cells were obtained. Also presented are the distributions of these probes in cells and simultaneous recording of changes in [Cl⁻]_i and pH_i during activation of Cl⁻-selective GlyR channels.

MATERIALS AND METHODS

PRODUCING AND CLONING OF ClopHensor VARIANTS

The ClopHensor, E²GFP-DsRedm, construct was mutated at two residues, H148G and V224L, of the GFP domain. Two sequential site-directed mutagenesis were performed using QuickChange II XL Site-Directed Mutagenesis Kit (Stratagene) following the manufacturer's protocol. Complementary primers were synthesized by Sigma-Aldrich with the following sequences: H148G-fw GAGTACAACACTACAACAGCGG CAACGTCTATATCATGG; H148G-rv CCATGATATAGACGTTGCCGCTGTTGTAGTTGTACTC; V224L-fw CTGCTGGAGTTCCTGAACGCGCCG; V224L-rv CGGCGGCGTTCAGGAACTCCAGCAG and were used to amplify the entire plasmid in a PCR reaction using high-fidelity polymerase. To eliminate template, the PCR reaction was digested with DpnI. The amplified mutated DNA was purified using Wizard SV Gel and the PCR Clean-up System kit (Promega), and transformed into *Escherichia coli* XL10-Gold ultracompetent cells (Novagen), which were then grown overnight on LB plates supplemented with 50 mg/l ampicillin at 37°C.

Four positive colonies were picked and grown overnight in 3 ml of LB-ampicillin at 37°C under shaking for mini prep DNA extraction (Wizard® Plus SV Minipreps DNA Purification;

Promega). All the constructs were verified by sequencing the entire insert. Finally, plasmids used for transfection were prepared using the QIAGEN Plasmid Plus Maxi kit.

EXPRESSION AND PURIFICATION OF THE H148G/V224L MUTANT

The recombinant GFP variant was expressed as Strep-tagged protein in *Escherichia coli* BL21 (Novagen) and harvested 20 h after induction with 1 mM IPTG at 30°C. Purification by affinity was carried out using Strep-Tactin Superflow 5-ml cartridges (IBA, GmbH, Germany), following the manufacturer's instructions, at 4°C in an AKTA Basic10 FPLC system (GE Healthcare Europe, Milan, Italy) with continuous monitoring of optical densities at 280 nm. The use of Cl⁻-free buffers in the final purification step ensured the complete removal of Cl⁻ from the preparation. Diethanolamine (DEA; 20 mM) in 50 mM K₂SO₄ adjusted to pH 8.5 was supplemented with a cocktail of protease inhibitors (Roche). Lysis was performed with an Ultrasonic Processor (Cole Parmer) (10 cycles of 30 s, output 6W, and 10 s cooling). FPLC was performed on an AKTA Basic10 FPLC system (GE Healthcare) using a Strep-Tactin Superflow 5 ml column (IBA Technology). Filtered lysate was loaded without a loop, and the flow rate was set at 3 ml/min. Elution of bound strep-tagged GFP was obtained in 75% washing buffer 2.5 mM desthiobiotin. Protein Concentration was determined by BCA™ Protein Assay (Pierce Protein Biology Products, USA).

CELL CULTURE PREPARATION

For fluorescence analysis of [Cl⁻]_i and [H⁺]_i, and also for immunocytochemistry and electrophysiology, ClopHensor constructs and human α1 GlyR subunits were expressed in CHO cells and in neurons of primary culture by means of Lipofectamin transfection.

Chinese hamster ovary (CHO-K1) cells were obtained from the American Type Tissue Culture Collection (ATCC, Molsheim, France) and maintained as previously described (Medina et al., 2000; Waseem et al., 2010). One day before the transfection, cells were plated onto coverslips (12–14 mm in diameter), which were placed inside 35-mm cell culture dishes with 2 ml of medium. CHO-K1 cells were transfected with approximately 1 μg/1 μl cDNA of constructs, using the Lipofectamine 2000 transfection protocol (Life Technology, USA). Three hours after the initial exposure of the cells to the cDNAs, a fresh cDNA-containing solution replaced the old one.

Neurons of hippocampal culture from 18-day rat embryos were dissociated using trypsin and plated at a density of 70,000 cells cm⁻² in minimal essential medium (MEM) supplemented with 10% NU serum (BD Biosciences, Le Pont de Claix, France), 0.45% glucose, 1 mM sodium pyruvate, 2 mM glutamine, and 10 IU ml⁻¹ penicillin–streptomycin as previously described (Buerli et al., 2007). On days 7, 10, and 13 of culture incubation, half of the medium was changed to MEM with 2% B27 supplement (Invitrogen).

Transfections of neuronal cultures at 7–10 days *in vitro* (DIV) were performed as described previously (Buerli et al., 2007; Pellegrino et al., 2011). Cells were used in experiments 2–5 days after transfection.

INTRACELLULAR Cl⁻ CALIBRATION OF ClopHensor VARIANTS

For Cl⁻ calibration we used whole-cell recordings with different Cl⁻ concentrations in recording pipettes. Whole-cell patch-clamp recordings on CHO cells were conducted 24–48 h after transfection, using an EPC-9 amplifier (HEKA Elektronik, Germany) at a holding potential -20 or -30 mV and at a temperature of 32–33°C. Cells were bathed in a solution containing (mM): NaCl 126; KCl 3.5; CaCl₂ 2; MgCl₂ 1.3; NaH₂PO₄ 1.2; NaHCO₃ 25; and D-glucose 10; equilibrated at pH 7.4 with 95% O₂ and 5% CO₂; 320 mOsm. The patch pipette solution contained (mM): KCl (0–135) or K-gluconate (0–135); MgCl₂ 2; MgATP 2; HEPES/KOH 10; and BAPTA 1; pH 7.3, 300 mOsm. A combination of K-gluconate and KCl at a constant K⁺ concentration of 135 mM were used for Cl⁻ calibration of ClopHensor constructs with six different Cl⁻ concentrations in the pipette solution ([Cl⁻]_p): 4, 10, 20, 60, 100, and 135 mM. Calibration curves were obtained by recording from 5–7 cells for each [Cl⁻]_p. The effectiveness of the cell dialysis and [Cl⁻]_i established after whole-cell penetration were checked by measurement of the reversal potential for glycine-induced currents. Glycine was applied locally using Picospritzer II (General Valve Corporation, USA) with a pipette positioned close to the soma of the recorded cell at different holding potentials.

All reagents were obtained from Sigma unless otherwise specified.

INTRACELLULAR pH CALIBRATION OF ClopHensor VARIANTS

For pH calibration b-escin permeabilization method was used (Waseem et al., 2010). In more detail, a range of HEPES-based extracellular solutions (150 mM K-Gluconate, 20 mM HEPES, and 10 mM D-glucose) with different pH values was created by adding HEPES powder for acidification of the solution or 1 M NaOH for alkalization.

To increase the permeability of the cell membrane to ions, 80 μM β-escin (Sigma, St Louis, USA) was added to the bath solution. β-escin was dissolved in water and prepared freshly for each experiment. This suspension was stable for about 2 h. The coverslip with cultured cells was placed into the recording chamber and superfused with escin-containing bath solution until cells become swollen, indicating the dissipation of ion gradients and coupled membrane potential in treated cells. The perfusing solution was then switched to the escin-free bath solution in order to avoid lysis of the cells. Thereafter, the fluorescence responses of ClopHensor constructs corresponding to specified H⁺ concentrations inside the cell were registered.

REAL-TIME FLUORESCENCE IMAGING

Fluorescence images were acquired using a customized digital imaging microscope. Excitation of cells at various wavelengths was achieved using a Polychrome V monochromator (Till Photonics, Germany). Light intensity was attenuated using neutral density filters. Emission wavelengths were controlled using a Lambda 10-3 controller (Sutter Instrument Company, USA). Fluorescence was visualized using an upright microscope Axioskop (Zeiss, Germany) equipped with a 60× water-immersion objective (n.a. 0.9; LumPlanFL, Olympus, USA). Fluorescent emitted light passed to a 16-bit digital camera

Andor iXon EM+ (Andor Technology PLC, Northern Ireland). Images were acquired on a computer via a DMA serial transfer. All peripheral hardware control, image acquisition and image processing were achieved using Andor iQ software (Andor Technology PLC). The average fluorescence intensity of each region of interest (ROI) was measured.

Cells expressing ClopHensor variants were excited at three wavelengths: 458 and 488 nm for Cl⁻/pH-sensitive E²GFP excitation, and 545 nm for excitation of DsRed-monomer. Fluorescent signals were recorded using a dual-band GFP/DsRed 493/574 dichroic mirror (Semrock Inc., USA) and two emission filters: 535 ± 15 nm for E²GFP emission and 632 ± 30 nm for emission of DsRed-monomer (both Chroma Technology Corporation, USA). The emission filters were mounted into the Lambda 10⁻³ Filter wheel (Sutter Instruments Company, Novato, USA).

The duration of excitation at each wavelength was usually 20 ms. The sampling interval was 5 s for the slow [Cl⁻]_i changes during the transition of the recorded cell in whole-cell configuration and was switched to 1 s for the fast [Cl⁻]_i transients in response to glycine application after establishing [Cl⁻]_i/[Cl⁻]_p equilibrium.

IMMUNOCYTOCHEMISTRY

For immunodetection of constructs in CHO cells or hippocampal neurons, cells in culture on coverslips expressing ClopHensor or its derivatives were fixed in PFA 4% (wt/vol) in 0.1 M phosphate buffer (PBS), pH 7.4, at room temperature for 15 min. After fixation, the cells were rinsed three times in PBS. In each experimental procedure, control and experimental cells were processed together to eliminate potential bias due to inherent variations in the intensity of the immunohistochemical labeling. For further staining, neurons were pre-incubated in 4% (vol/vol) normal goat serum in PBS containing 0.5% Triton X-100 (vol/vol) for 1 h at room temperature to block secondary-antibody binding sites. Coverslips with neurons expressing ClopHensor were then incubated overnight at 4°C with rabbit anti-GFP polyclonal antibody (Invitrogen) diluted 1:4000 in PBS with 4% normal goat serum and mouse monoclonal anti-MAP2 antibodies (1:2000). Incubations with the primary antibodies were performed at room temperature and slow shaking for 1 h, then overnight at 4°C. Samples were rinsed three times in PBS. As secondary antibodies, either a Cy3-conjugated anti-rabbit IgG for detection of GFP or a Cy5-conjugated anti-mouse IgG for detection of MAP2 (all 1:1500; Molecular Probes, Invitrogen) were used. Samples were then rinsed twice in PBS with one additional washing in PBS containing Hoechst 33342 (10 mg/ml; Sigma-Aldrich). Coverslips were mounted using Fluoromount™ (Sigma-Aldrich, St Louis, MO, USA).

Images were acquired using a Leica SP5C confocal microscope using 40–60× oil-immersion objectives. Micrographs shown here are digital composites of Z-series scans of 5–15 optical sections through a depth of 1–6 μm. Final images were constructed with ImageJ software.

In vitro ANALYSIS

In vitro characterization of H⁺ and Cl⁻-binding properties was performed by analysing fluorescence spectra variations with a

multimodal plate reader (EnSpire; PerkinElmer) in 96-well plates. Acetic acid buffer (50 mM) was used to adjust the solution pH to around 5.2, 100 mM PBS was used for the pH range 5.2–8.0 and 20 mM DEA buffer was used for the pH range 8.0–8.5. Temperature was kept constant at 20 ± 0.5°C and fluorescence spectra were acquired with 1-nm steps. Protein concentration was always kept constant at about 1 μM throughout every pH and Cl⁻ titration.

The proton dissociation constant (pK_a) was obtained by fitting fluorescence data with the equation:

$$F = \frac{A_1 + A_2 10^{(pK_a - pH)}}{1 + 10^{(pK_a - pH)}}$$

which describes the effect of environmental proton concentration (pH) on chromophore fluorescence (F).

The chloride dissociation constant (K_d^{Cl}) was obtained by fitting data with a 1:1 Langmuir binding model (Arosio et al., 2007):

$$S = \frac{S_0 + S_1 \cdot [Cl] / K_d^{Cl}}{1 + [Cl] / K_d^{Cl}}$$

RESULTS

DESCRIPTION OF THE CONSTRUCTS AND *in vitro* ANALYSIS

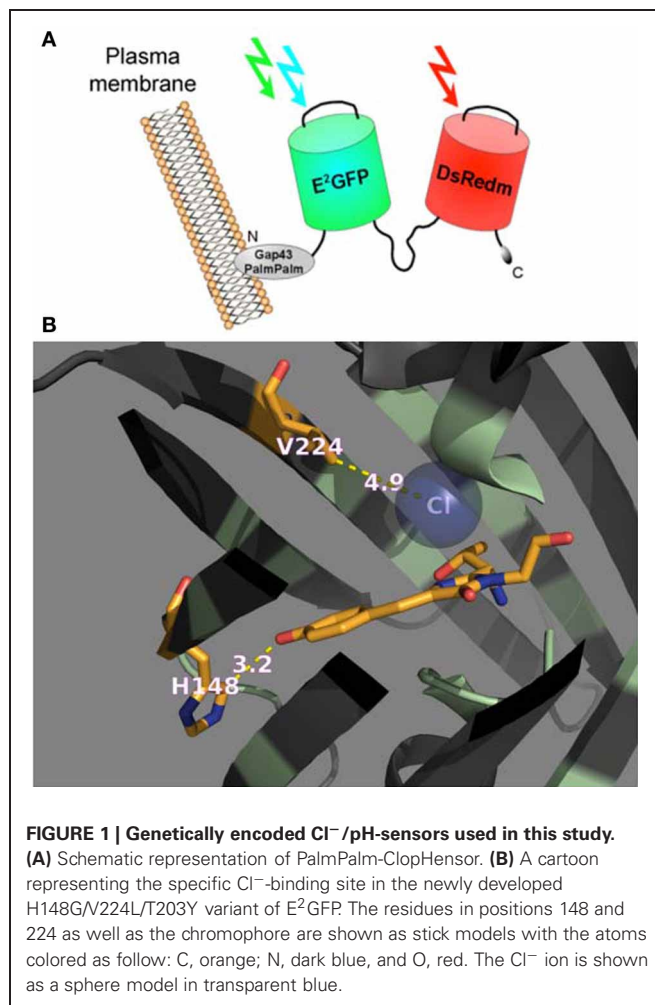
In this study we performed calibration and functional characterization of the original ClopHensor and doubly palmitoylated membrane-targeted constructs (Arosio et al., 2010), as well as a new ClopHensor variant with mutations designed to increase Cl⁻ affinity and shift pK_a to more alkaline values (Figure 1B).

ClopHensor was designed by linking two fluorescent proteins, E²GFP and DsRed-monomer by means of a flexible 20-amino-acid spacer. To target the sensor to the inner interface of the plasma membrane, we inserted the N-terminal 20 residues of GAP-43, which contain two palmitoylation sites, at the ClopHensor N-terminus (Figure 1A). DsRed fluorescence, which is not affected by pH or Cl⁻ changes, was shown to provide an excellent normalization signal leading to ratiometric Cl⁻ sensing, free from the influence of sensor concentration in living neurons.

Following purification of the E²GFP-H148G-V224L mutant, Cl⁻-binding properties of the recombinant protein were investigated *in vitro* by measuring fluorescence spectra at constant pH and increasing Cl⁻ concentration, from 0 to 310 mM (Figures 2A,B). Because of the cooperative interaction between Cl⁻ and H⁺ binding, Cl⁻ titrations were measured at various pH values (from 5.25 to 8.9) and K_d values were analyzed with an infinite cooperative model (Arosio et al., 2007; Bregestovski and Arosio, 2011) with two interacting binding sites, one for Cl⁻ and one for H⁺:

$$K_d = \frac{K_d^0 \cdot (1 + 10^{(pK_a - pH)})}{10^{(pK_a - pH)}}$$

containing two fitting parameters for the probe: pK_a and K_d⁰, the Cl⁻ dissociation constant for the fully protonated form (present at pH ≤ 5).



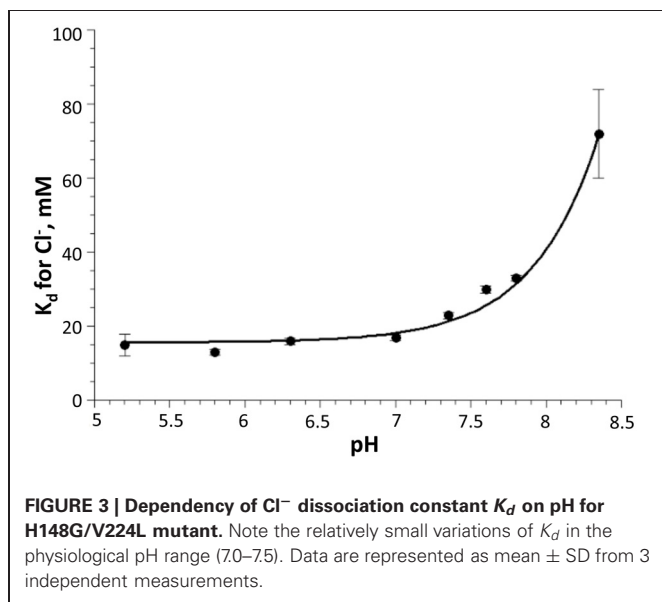
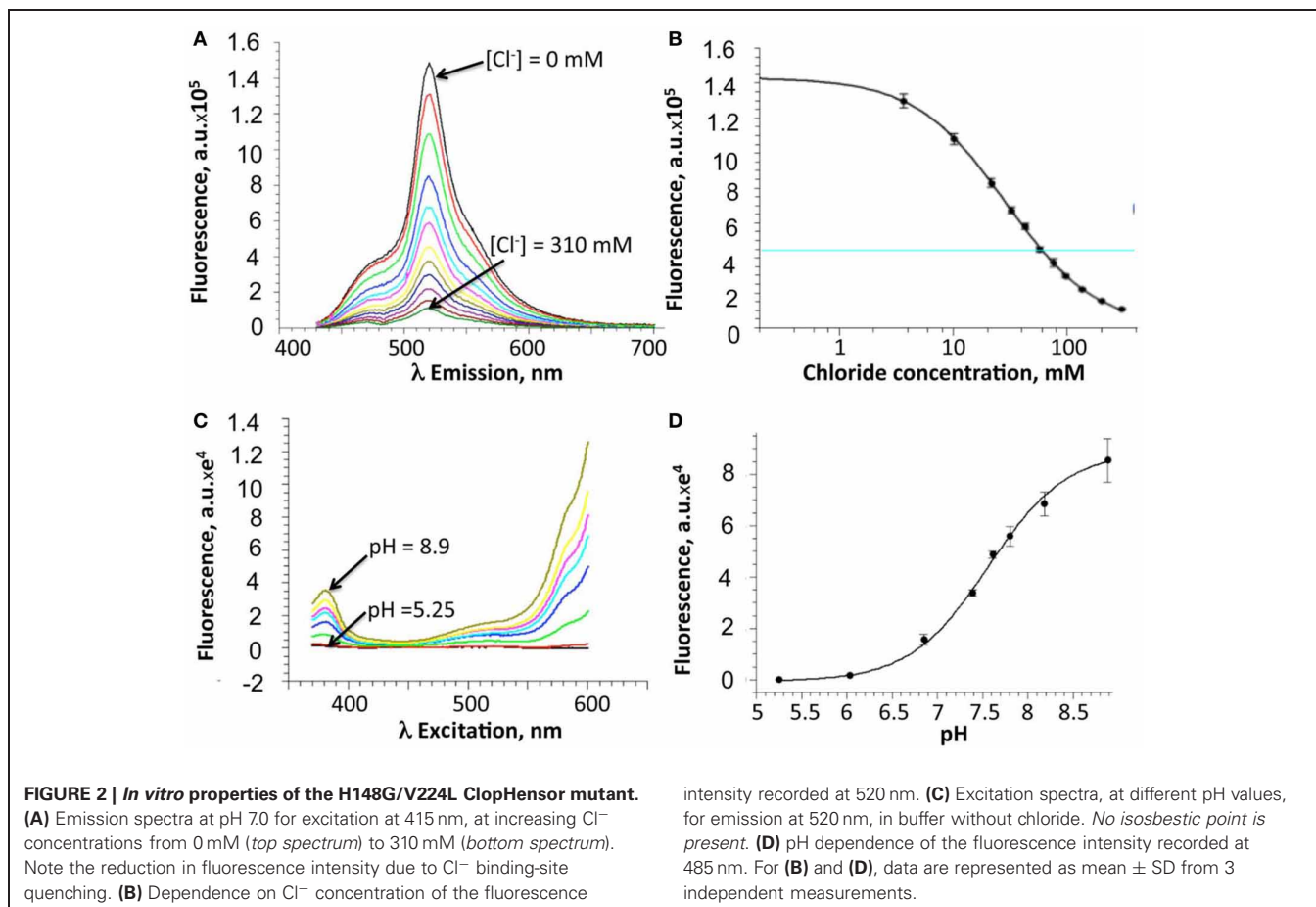
The proton binding equilibrium was studied and the pK_a value was determined from spectra acquired at different pH buffer conditions, ranging from 5.25 to 8.9, in the absence of halogens (Figures 2C,D).

Precisely, the Cl⁻ and H⁺ binding thermodynamic parameters for H148G/V224L were found to be pK_a = 7.9 ± 0.05 (best-fit ± SD) and K_d⁰ = 19 ± 1.5 mM.

Overall, *in vitro* analysis at various pH and Cl⁻ conditions revealed that the H148G/V224L mutant is suited for Cl⁻ concentration measurements with reduced pH dependence in the physiological range. Indeed, in the pH range between 6.8 and 7.8 K_d changes were between 18 and 30 mM (Figure 3).

DISTRIBUTION OF THE CONSTRUCTS IN CELLS

To determine the intracellular distribution of ClopHensor constructs, we transiently expressed them in CHO cells and in neurons of dissociated hippocampal culture (see Materials and Methods). Expression was already observable 12 h after transfection and remained at high levels for at least 4 days. The fluorescence of the ClopHensor was distributed homogeneously throughout CHO cells (Figures 4A,B, left) and in neurons (Figure 4C). The fluorescence was bright with no preferential



staining of membrane or intracellular organelles, as illustrated by the fluorescence profile (Figure 4B, left).

Distribution of the H148G/V224L mutant was also cytoplasmic, similar to those for ClopHensor (Figures 4A,B, middle).

In contrast, the PalmPalm-ClopHensor typically showed near-membrane distribution; however, a small fraction of the probe could still be observed in the cytoplasm (Figures 4A,B, right).

These observations indicate that ClopHensor and H148G/V224L mutants exhibit cytoplasmic intracellular distributions while the PalmPalm-ClopHensor construct is typically localized to the plasma membrane and the Golgi region, as it was shown previously (McCabe and Berthiaume, 1999).

INTRACELLULAR CALIBRATION OF THE CONSTRUCTS

Calibration for Cl⁻

In order to evaluate the dynamic range and sensitivity of constructs to ions, we co-expressed them in CHO cells with Cl⁻-selective GlyR channels and performed simultaneous monitoring of whole-cell currents and fluorescent signals. Whole-cell recordings were performed with different Cl⁻ concentrations in the pipette solution ($[Cl^-]_p = 4\text{--}135\text{ mM}$). To activate GlyR channels, glycine was applied using pressure pulses. For this, a pipette containing 200 μM glycine dissolved in extracellular solution was advanced to within 30–50 μm of the recorded cell (Figure 5A).

For ratiometric monitoring, following the approach presented in Arosio et al. (2010) we used the ratio $R_{Cl} = F_{458}/F_{545}$ for measurement of intracellular Cl⁻ concentration ($[Cl^-]_i$), and for calculating intracellular pH values (pH_i) the ratio $R_{pH} = F_{488}/F_{458}$ was used.

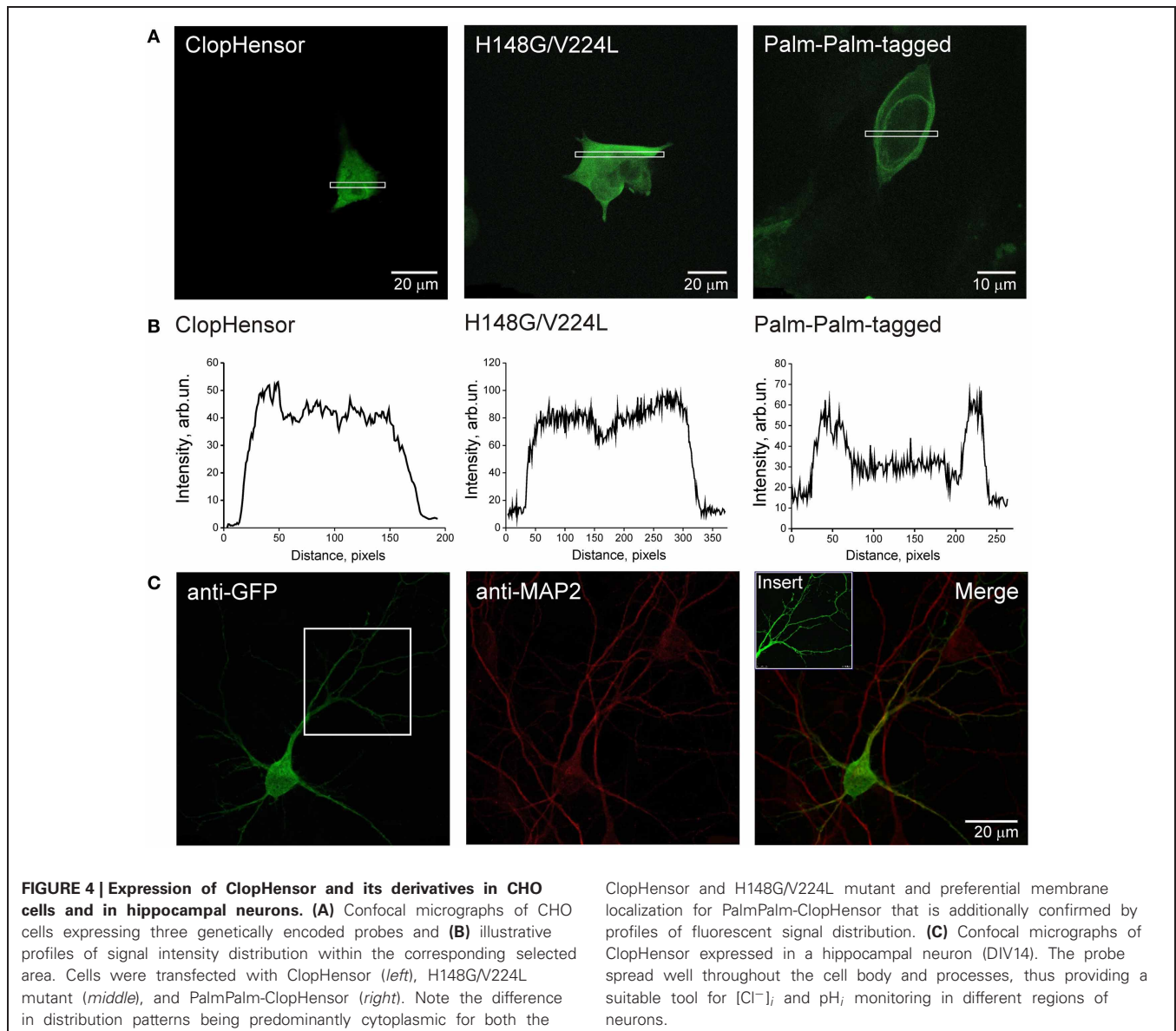


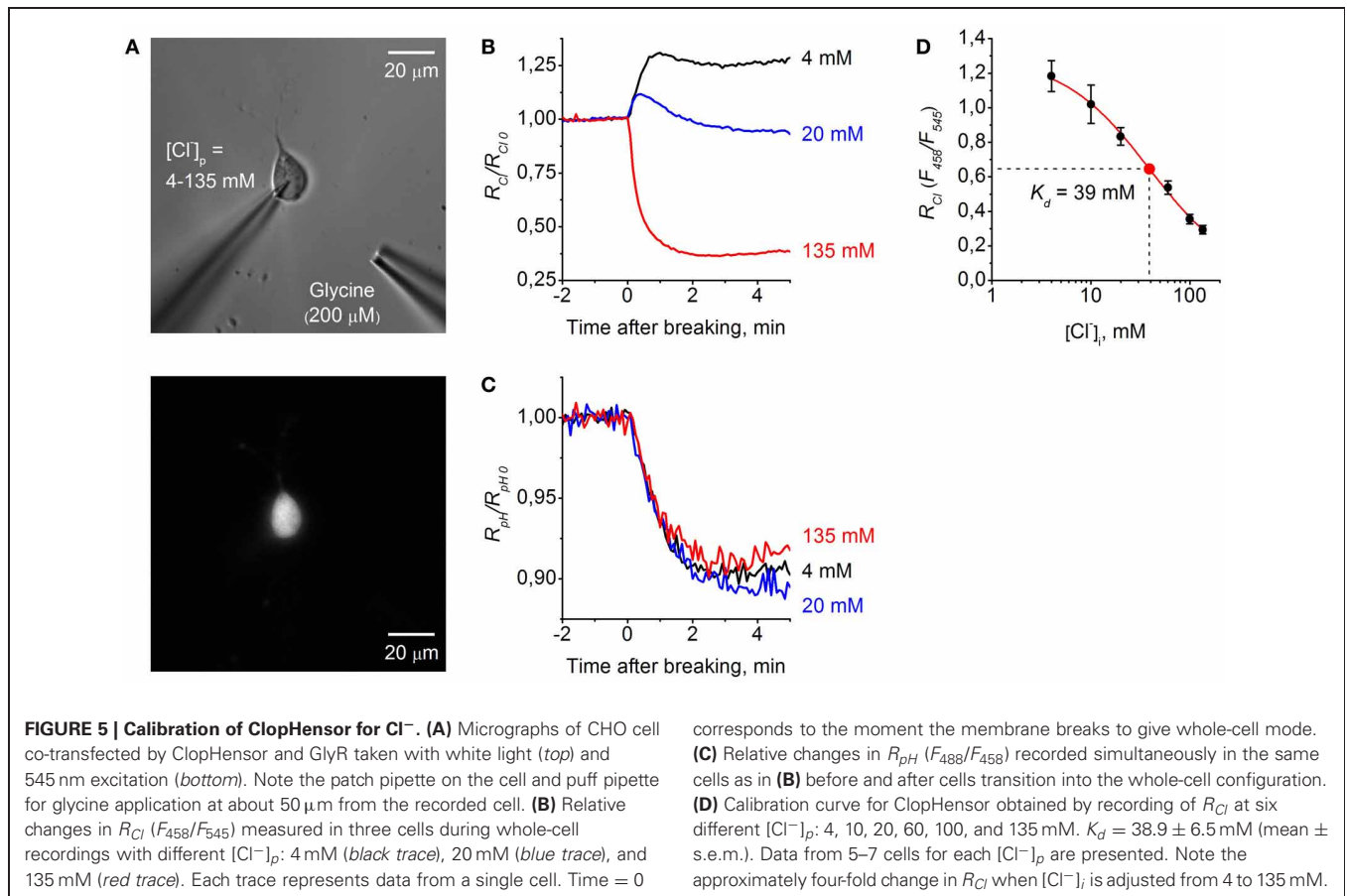
Figure 5B illustrates relative changes in [Cl⁻]_i during whole-cell recordings from three cells, using pipettes with solutions containing different Cl⁻ concentrations. The graph represents changes in R_{Cl}/R_{Cl_0} with time, where R_{Cl_0} corresponds to [Cl⁻]_i in the cell-attached mode, i.e., to the native concentration of Cl⁻ in the cytoplasm of CHO cells. Obtaining the whole-cell configuration by rupturing the membrane with a pipette containing 135 mM Cl⁻ resulted in a strong decrease in R_{Cl}/R_{Cl_0} , corresponding to elevation of [Cl⁻]_i. In contrast, on rupture of the membrane with the pipette containing 4 mM Cl⁻ an increase in R_{Cl}/R_{Cl_0} was observed. Transition to the whole-cell configuration with a pipette containing 20 mM Cl⁻ produced a short transient, increasing with stabilization of R_{Cl}/R_{Cl_0} at a level close to those in the cell-attached configuration. This reflects the fact that in the recorded cell the native value of [Cl⁻]_i in cytoplasm is close to the [Cl⁻]_p of 20 mM.

Simultaneous monitoring of relative pH changes (R_{pH}/R_{pH_0}) showed that after transition to the whole-cell configuration pH_i similarly decreased for all three cells recorded with different [Cl⁻]_p but with identical pH (**Figure 5C**).

Calibration curves for Cl⁻ obtained from six different [Cl⁻]_p in physiological conditions were best fitted by a Logistic Dose-Response Sigmoidal Fit using the OriginPro 8.5 program with the formula:

$$R_{Cl} = A2 + \frac{A1 - A2}{1 + \left(\frac{[Cl^-]_i}{K_d}\right)^p}$$

where R_{Cl} is the fluorescence ratio for Cl⁻ (F_{458}/F_{545}), K_d is the dissociation constant for Cl⁻ binding, A1 and A2 are, respectively, the minimum and maximum asymptotic values of R_{Cl} , and p is the power value.



From this formula we obtained the equation for $[Cl^-]_i$ values recalculation:

$$[Cl^-]_i = K_d \cdot \left(\frac{A1 - A2}{R_{Cl} - A2} - 1 \right)^{\frac{1}{p}}$$

For ClopHensor the values of constants obtained from fitting the curve were the following: $K_d = 38.9 \pm 6.5$ mM, $A1 = 1.29$, $A2 = 0.01$, and $p = 1$ (Figure 5D).

Similar analyses were performed for PalmPalm-ClopHensor and H148G/V224L mutants (Figure 6). They demonstrated that changes in R_{Cl}/R_{Cl0} observed at transitions from cell-attached to whole-cell configuration in cells expressing PalmPalm-ClopHensor were similar to those for ClopHensor (Figure 6A). From the calibration curve for PalmPalm-ClopHensor the following fitting parameters were obtained: $K_d = 46.8 \pm 3.8$ mM, $A1 = 1.15$, $A2 = 0.29$, and $p = 1.02$ (Figure 6B). While this value of K_d was higher than for ClopHensor, the difference was non-significant ($P > 0.05$, t -Student).

For cells expressing the H148G/V224L mutant, R_{Cl}/R_{Cl0} when recording with the pipette containing 20 mM of Cl⁻ was almost midway between values recorded with pipettes containing 135 and 4 mM (Figure 6C), suggesting higher sensitivity of the construct to Cl⁻. This was confirmed by obtaining the calibration curve for the H148G/V224L mutant. The constants were the following: $K_d = 21.4 \pm 4.8$ mM, $A1 = 1.58$, $A2 = 1.06$, and $p =$

1.51 (Figure 6D). Statistical analysis of K_d values for ClopHensor vs. H148G/V224L showed that they were significantly different ($P < 0.05$, t -Student) and for PalmPalm vs. H148G/V224L K_d values were also significantly different ($P < 0.01$, t -Student). The dynamic range of R_{Cl} changes for the mutant was, however, smaller than those for the ClopHensor, which reduces the benefits arising from the increased affinity.

Calibration for pH

As the H148G/V224L mutation caused a shift in the pK_a , we performed comparative pH calibrations of ClopHensor and H148G/V224L mutant. For pH calibration, the β -escin method was used (see Materials and Methods).

Sequential exchanging of bath solutions with different pH values produced corresponding shifts in the fluorescence pH ratio (Figure 7).

For pH calibration curves a Dose-Response Sigmoidal Fit (OriginPro 8.5) with the following formula was used:

$$R_{pH} = B1 + \frac{B2 - B1}{1 + 10^{p(pK_a - pH_i)}}$$

where R_{pH} is the fluorescence ratio for pH (F_{488}/F_{545}), pK_a is the acid dissociation constant for H⁺ binding, B1 and B2 are, respectively, the minimum and maximum asymptotic values of R_{pH} , and p is the power value.

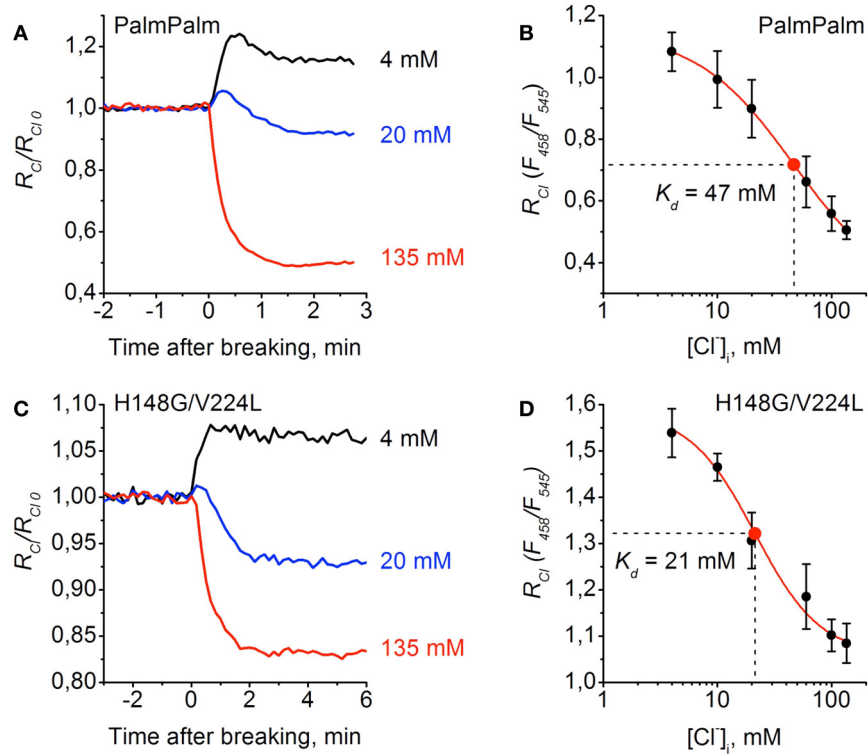


FIGURE 6 | Calibration of PalmPalm-ClopHensor and H148G/V224L mutant for Cl⁻. (A) and (C) Relative changes in R_{Cl} (F_{458}/F_{545}) of PalmPalm-ClopHensor (A) and the H148G/V224L mutant (C) measured in three cells during whole-cell recordings with different $[Cl^-]_p$: 4 mM (black trace), 20 mM (blue trace), and 135 mM (red trace). Each trace represents the data from a single cell. Time = 0 corresponds to the moment the membrane breaks to give whole-cell mode. (B) and (D) Calibration curve for

PalmPalm-ClopHensor (B) and H148G/V224L mutant (D) obtained by recording R_{Cl} at six different $[Cl^-]_p$: 4, 10, 20, 60, 100, and 135 mM. For PalmPalm-ClopHensor $K_d = 46.8 \pm 3.8$ mM and for H148G/V224L mutant $K_d = 21.4 \pm 4.8$ mM. Data from 5–7 cells for each $[Cl^-]_p$ are presented. Note the approximately two-fold changes in R_{Cl} for PalmPalm-ClopHensor when $[Cl^-]_i$ changed from 4 to 135 mM and the much smaller dynamic range of R_{Cl} for H148G/V224L mutant.

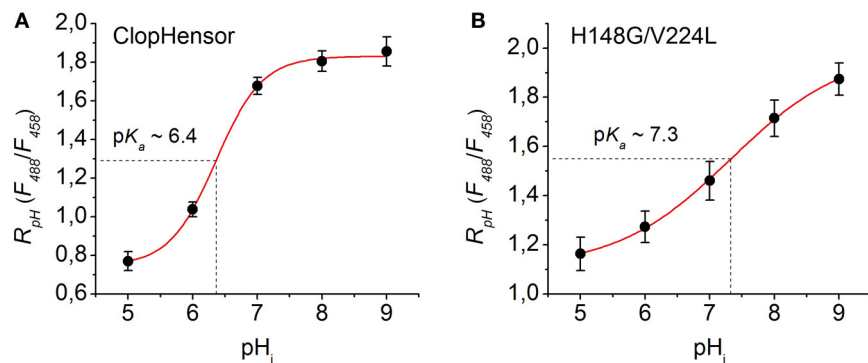


FIGURE 7 | Calibration of ClopHensor and H148G/V224L mutant for pH. (A) and (B) Changes in R_{pH} (F_{488}/F_{458}) of ClopHensor (A) and H148G/V224L mutant (B) measured in transfected partially permeabilized CHO cells perfused with bath medium of different pH values (see Materials and Methods for details).

From this formula we obtained the equation for pH_i values recalculation:

$$pH_i = pK_a - \frac{1}{p} \cdot \log \left(\frac{B2 - B1}{R_{pH} - B1} - 1 \right)$$

Based on the fitting of the experimental curves the constants values obtained were: $pK_a = 6.36$, $B1 = 0.75$, $B2 = 1.83$, and $p = 1.21$ for ClopHensor (Figure 7A) and $pK_a = 7.32$, $B1 = 1.11$, $B2 = 1.99$, and $p = 0.49$ for the H148G/V224L mutant (Figure 7B). In comparison with ClopHensor, the dynamic range

of R_{pH} changes and the slope of the calibration curve characterizing by the constant p for the mutant were smaller, indicating its lower pH sensitivity.

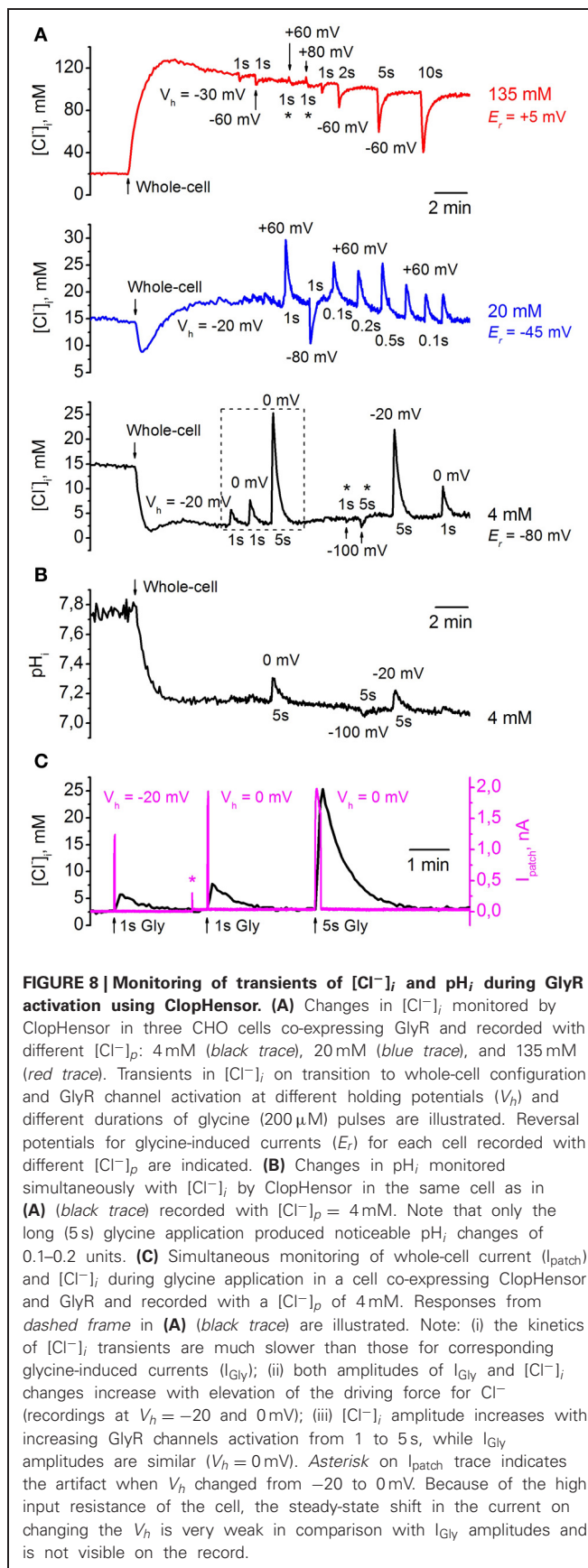
Transients of $[\text{Cl}^-]_i$ and pH_i during activation glycine receptor channels

We further performed comparative analysis of the three variants of ClopHensor while monitoring $[\text{Cl}^-]_i$ and pH_i transients induced by activation of Cl^- -selective GlyR channels in CHO cells.

Figure 8A illustrates changes in $[\text{Cl}^-]_i$ monitored from three CHO cells co-expressing ClopHensor with GlyR and recorded with different $[\text{Cl}^-]_p$: 4 mM (black trace), 20 mM (blue trace), and 135 mM (red trace). Values of $[\text{Cl}^-]_i$ recalculated from R_{Cl} (F_{458}/F_{545}) were obtained using the calibration curve in **Figure 5D**. The transition to whole-cell configuration (holding potential $V_h = -30$ mV) with a pipette containing 135 mM Cl^- caused a strong elevation of $[\text{Cl}^-]_i$. After peaking, $[\text{Cl}^-]_i$ slowly declined, presumably due to pumping out of Cl^- by transporters (see, for instance, Pellegrino et al., 2011) or through weakly activated Cl^- -selective ionic channels (Friedel et al., 2013). Application of 200 μM glycine to the cell induced a transient decrease in $[\text{Cl}^-]_i$, the amplitude of which depended on the duration of the applied pulses of the agonist and values of V_h . The holding potential was transiently changed to the values indicated on the figure for each pulse of glycine. Application of the agonist for 1 s caused a decrease of 5–10 mM, while activation of GlyRs for 10 s resulted in decrease of about 60 mM (**Figure 8A**, red trace). After transiently diminishing, $[\text{Cl}^-]_i$ recovered to its initial steady-state level. Interestingly, for the cell recorded with $[\text{Cl}^-]_p = 135$ mM it was difficult to increase $[\text{Cl}^-]_i$ in conditions under which the glycine-induced current caused a strong influx of Cl^- , i.e., at $V_h = +60$ and even at $+80$ mV (**Figure 8A**, asterisks, red trace). In this cell the reversal potential (E_r) for I_{Gly} was about +5 mV, as predicted for symmetric intracellular-extracellular Cl^- concentrations.

Simultaneous recording of $[\text{Cl}^-]_i$ and ionic currents (**Figure 8C**) showed two main features. First, the kinetics of $[\text{Cl}^-]_i$ transients were much slower than those for glycine-induced currents. This is in accordance with previous observations from monitoring of Cl^- transients using MQAE-mediated fluorescence (Marandi et al., 2002) or BioSensor-GlyR (Mukhtarov et al., 2008). Second, $[\text{Cl}^-]_i$ amplitude increased both on elevation of the driving force for Cl^- and on prolongation of glycine applications. As illustrated in **Figure 8C**, the amplitude of $[\text{Cl}^-]_i$ transients induced by 1-s pulses of glycine increased in parallel with GlyR current amplitude at changes of V_h from -20 to 0 mV. At constant V_h (0 mV), prolongation of glycine pulses from 1 to 5 s caused an increase in $[\text{Cl}^-]_i$ amplitude although an increase in GlyR current amplitude was not observed.

When transition to whole-cell configuration ($V_h = -20$ mV) was produced with a pipette containing 20 mM Cl^- a brief transient decrease in $[\text{Cl}^-]_i$ with subsequent stabilization at a level close to that in the cell-attached configuration was observed (**Figure 8A**, blue trace). Even at short (100 ms) glycine applications, substantial transients in $[\text{Cl}^-]_i$, of about 5 mM, were



recorded with $V_h = +60$ mV; these reversed in direction when V_h was changed from $+60$ to -80 mV. E_r for glycine-induced currents in this cell was about -45 mV (Figure 8A, blue trace).

With a pipette containing 4 mM Cl^- , transition to whole-cell configuration ($V_h = -20$ mV), as predicted, caused a decrease in $[\text{Cl}^-]_i$ (Figure 8A, black trace). At $V_h = -20$ and 0 mV, application of 200 μM glycine induced strong $[\text{Cl}^-]_i$ transients whose amplitudes increased with prolongation of pulse duration; these increases were about 3–5 and 18–22 mM for glycine pulses of 1 and 5 s, respectively. Similarly to high $[\text{Cl}^-]_p$, for the cell recorded with a $[\text{Cl}^-]_p$ of 4 mM it was difficult to decrease $[\text{Cl}^-]_i$ in conditions under which glycine-induced current caused efflux of Cl^- , i.e., at $V_h = -100$ mV (Figure 8A, asterisks, black trace). E_r for glycine-induced currents in this cell was about -80 mV.

Simultaneous monitoring of pH_i in the same cell as that shown in Figure 8A, black trace, demonstrated a decrease in pH_i of about 0.6 units after transition to whole-cell configuration (Figure 8B). Values of pH_i recalculated from R_{pH} (F_{488}/F_{458}) were obtained using the calibration curve in Figure 7A. The changes in $[\text{Cl}^-]_i$ up to 5 mM (with 1-s glycine applications) did not produce marked changes in pH_i , while the changes in $[\text{Cl}^-]_i$ to ≥ 20 mM (with 5-s glycine applications) caused changes in pH_i of 0.1–0.2 units (Figure 8B).

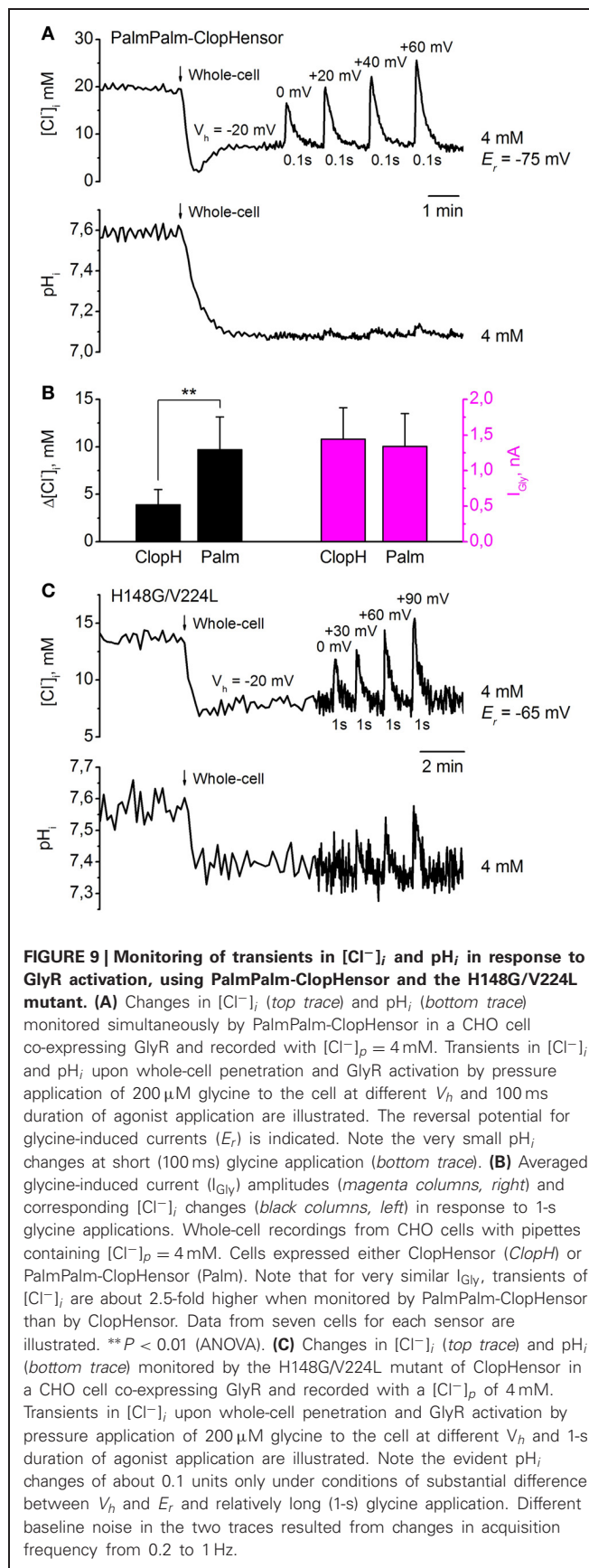
These observations indicate that ClopHensor is a reliable tool for long-lasting simultaneous monitoring of intracellular Cl^- and pH in living cells.

Next we analysed the properties of ClopHensor derivatives: the membrane-targeted PalmPalm-ClopHensor and the H148G/V224L ClopHensor mutant.

Figure 9A illustrates simultaneous monitoring of $[\text{Cl}^-]_i$ (top trace) and pH_i (bottom trace) by PalmPalm-ClopHensor in CHO cells co-expressing GlyR and recorded with a $[\text{Cl}^-]_p$ of 4 mM. Values of $[\text{Cl}^-]_i$ recalculated from R_{Cl} (F_{545}/F_{458}) were obtained using the calibration curve in Figure 6B. The transition to whole-cell configuration ($V_h = -20$ mV) produced a decrease in $[\text{Cl}^-]_i$ similar to those monitored by ClopHensor. Short (100 ms) application of 200 μM glycine to the cell induced transient increases in $[\text{Cl}^-]_i$ that were dependent on the difference between the holding potential and the reversal potential of glycine-induced currents.

Transients in $[\text{Cl}^-]_i$ varied from about 10 mM at $V_h = 0$ mV to about 20 mM at $V_h = +60$ mV (Figure 9A, top trace). E_r for glycine-induced currents in this cell was about -75 mV. These changes were much higher than would be found when monitoring with ClopHensor at similar stimuli. This was confirmed on analysis of $[\text{Cl}^-]_i$ transients when recording from cells expressing these two sensors. As illustrated in Figure 9B, very similar GlyR currents induced $[\text{Cl}^-]_i$ transients whose amplitude appeared about 2.5-fold higher when monitored by PalmPalm-ClopHensor.

As the molecular organization of PalmPalm-ClopHensor is identical to that of ClopHensor, the recalculation of pH_i values from R_{pH} (F_{488}/F_{458}) was done using the calibration curve in Figure 7A. Similarly, after transition to whole-cell configuration, pH_i decreased by about 0.5 units (Figure 9A, bottom trace). Very weak pH_i transients were observed (Figure 9A, bottom trace) with short (100 ms) glycine applications even at $V_h = +60$ mV, while the transient increase in $[\text{Cl}^-]_i$ in this cell was about 20 mM (Figure 9A, top trace).



When the H148G/V224L mutant of ClopHensor was co-expressed with GlyR in CHO cells recorded with $[Cl^-]_p = 4$ mM, a decrease in $[Cl^-]_i$ on transition to whole-cell configuration ($V_h = -20$ mV) was also observed (**Figure 9C**, top trace). Values of $[Cl^-]_i$ recalculated from $R_{Cl}(F_{545}/F_{458})$ were obtained using the calibration curve in **Figure 6D**. Again, glycine-induced transient increases in $[Cl^-]_i$ depended on the difference between V_h and E_r . Transients in $[Cl^-]_i$ with 1-s glycine application varied from about 5 mM at $V_h = 0$ mV to about 10 mM at $V_h = +90$ mV. E_r for glycine-induced currents in this cell was about -65 mV.

Simultaneous monitoring of pH_i in the same cell reveal that the $[Cl^-]_i$ transients below 5 mM did not produce marked changes in pH_i . The recalculation of pH_i values from $R_{pH}(F_{488}/F_{458})$ was done using the calibration curve in **Figure 7B**. An evident pH_i increase (of about 0.1 units) was recorded only when there was a substantial difference between V_h and E_r and relatively long (1-s) glycine application, which induced a $[Cl^-]_i$ elevation of about 10 mM (**Figure 9C**, bottom trace).

These results demonstrate that PalmPalm-ClopHensor, which has preferable membrane targeting, could be of interest for near-membrane measurement of intracellular Cl⁻ and pH changes. Another construct, the H148G/V224L mutant of ClopHensor, with a higher affinity for Cl⁻, possesses the smallest dynamic range of $R_{Cl}(F_{458}/F_{545})$ changes and this could be a limiting factor in some cases due to lower signal/noise ratio.

DISCUSSION

Development of genetically encoded probes for non-invasive monitoring of ions and protein function has opened powerful routes for the analysis of a variety of physiological problems and functions of various cell types under different experimental conditions. These probes are non-toxic, capable of remaining stable in cells for a long time, can be expressed in specific cellular compartments and are suitable for production of transgenic models.

Here, we describe the calibration in living cells, the cytoplasmic distribution, and examples of simultaneous monitoring of intracellular Cl⁻ and H⁺ of three genetically encoded sensors: (1) ClopHensor; (2) its variant designed to have preferential membrane targeting due to the addition to the N-terminus of a short peptide containing two palmitoylation sites (PalmPalm-ClopHensor) (Arosio et al., 2010), and (3) ClopHensor containing mutations of E²GFP (H148G/V224L) aimed at increasing the Cl⁻ affinity of the sensor.

Following transient expression in CHO cells, ClopHensor and the H148G/V224L mutant exhibit cytoplasmic intracellular distribution while the PalmPalm-ClopHensor construct, as expected, is preferentially localized in the vicinity of membranes. Upon transfection of rat dissociated hippocampal cultures with cDNA of ClopHensor, the probe shows a strong cytoplasmically distributed fluorescence in the soma and neuronal processes. These observations indicate that all three sensors can be easily expressed in various cell types and detected in different, even very small, areas of cells.

Calibration analysis performed on CHO cells using patch pipettes containing different concentrations of Cl⁻, and also

β-escin permeabilization with bath medium having different pH (Waseem et al., 2010) revealed that the constructs exhibit different sensitivity to Cl⁻ and H⁺. While ClopHensor and PalmPalm-ClopHensor probes demonstrated K_d for Cl⁻ of about 40 mM, for the H148G/V224L mutant this value was about 20 mM, indicating its higher affinity. However, the mutant exhibits a smaller dynamic range of R_{Cl} , which could be a limiting factor for monitoring Cl⁻ in cells containing low $[Cl^-]_i$ or with small changes in Cl⁻ concentration. Poor dynamic range could be, at least partially, explained by different spectral properties of H148G/V224L mutant (**Figures 2A,C**) from those for ClopHensor (Arosio et al., 2010). In the present study, for all constructs including the H148G/V224L mutant, the calibration measurements were performed using the same excitation wavelengths, GFP/DsRed dichroic mirror and emission filters (see Materials and Methods). We suggest that this range can be considerably extended in the future by better selection of excitation and emission parameters.

Calibration of pH in CHO cells using the β-escin method showed that the pK_a for the H148G/V224L mutant is strongly shifted to alkaline values. This is consistent with *in vitro* measurements: $pK_a = 7.9 \pm 0.05$. *In vitro* analysis also demonstrated that K_d for Cl⁻ of the H148G/V224L mutant exhibits relatively small pH dependency over a wide pH range: from about 18 mM at pH 6.5 to about 30 mM at pH 7.8. This suggests that the H148G/V224L mutant is a useful tool for $[Cl^-]_i$ measurements in experimental models with high pH variations.

Our previous observations demonstrated that when using the CFP/YFP-based Cl-Sensor, long or frequent acquisition causes strong “bleaching” of Cl⁻-sensitive YFP resulting in changes in fluorescence parameters during ratiometric measurement of $[Cl^-]_i$. This problem is discussed in detail in the paper by Friedel et al. (2013, this issue). In contrast, E²GFP-DsRedm-based ClopHensor exhibits remarkable stability (see, for instance, **Figures 8, 9**), providing an excellent tool for long-lasting reliable monitoring and with variable acquisition rate.

Our observations of the effects of activation of Cl⁻-selective GlyR channels on $[Cl^-]_i$ indicate that it is a highly dynamic parameter, which can be strongly changed by over-activation of Cl⁻-selective channels or activity of other proteins involved in regulation and determination of physiological $[Cl^-]_i$ in living systems. Indeed, activation of GlyRs for several seconds caused changes in $[Cl^-]_i$ of more than 10–20 mM (**Figures 8, 9**).

Together our experiments demonstrate that these three ClopHensor constructs are suitable tools for stable, long-lasting, non-invasive monitoring of $[Cl^-]_i$ and pH_i in different cell types.

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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