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A subcellularly targeted photocaged inhibitor for mitochondrial carbonic anhydrase V†

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We report the first photoactivatable inhibitor specifically targeting mitochondrial CA-V, a promising anti-obesity target. It features a sulfonamide metal-binding group caged with a photoremovable coumarin and guided to mitochondria via a pyridinium moiety. Light exposure removes the cage, enabling precise spatial and temporal inhibition of mitochondrial CA-V activity.

Carbonic anhydrase (CA) is a zinc-dependent metalloenzyme that catalyzes the reversible hydration of carbon dioxide and bicarbonate.¹ There are 15 known CA isoforms; of these, CA-II and CA-IX have been most extensively studied for inhibitor development.^{2,3} However, mitochondrial isoforms CA-VA and CA-VB, which show 66% sequence similarity, are relatively less explored.^{4,5} CA-VA is most abundant in the liver⁶ and is involved in crucial biosynthetic processes including ureagenesis, lipogenesis, and others.^{7–10} Dysregulation of CA-VA is linked to obesity, diabetes, hyperammonemia, growth deficiencies, and brain-related oxidative stress illnesses.^{6,11–17} CA-VB has a broad tissue distribution, however much less is known about its biological function.^{6,18} Despite the health relevance of CA-VA and CA-VB, limited research has been conducted compared to the more ubiquitous CA-II and CA-IX due to the challenge of selectively targeting CA-V isoforms within cells and organisms.

Current methods for studying CA-V isoforms can be categorized into three main approaches. First, knockout (KO) mouse models can provide valuable insights into the overall impact of complete CA-V deficiency in organisms.¹³ However KO models fall short in addressing the specific timing, mechanisms, and causality of CA-V dysregulation, particularly in relation to mitochondrial dysfunction. A second approach involves inhibiting CA-V within isolated mitochondria.^{8,19–21} While this method offers kinetic and biochemical data for

CA-V, it requires that mitochondria are extracted from cells, leaving the interactions of CA-V within the broader cellular context unexplored. Finally, CA-V-specific inhibitors have been pursued and tested extensively *in vitro*, some with CA-V selectivity ratios as high as 3800, though their activation cannot be controlled in a spatiotemporal manner.^{22–34}

In this study, we report a novel photocaged CA inhibitor (PhotoCaged Ethoxzolamide Mitochondria-targeted, **PCEM**) that can be selectively delivered to mitochondria without inhibiting cytosolic or other CA isoforms in the process (Fig. 1). This tool uses an aryl sulfonamide ($-\text{SO}_2\text{NH}_2$) for coordination to the Zn^{2+} in the CA active site. This ligand is blocked from interacting with CA by a photoremovable protecting group

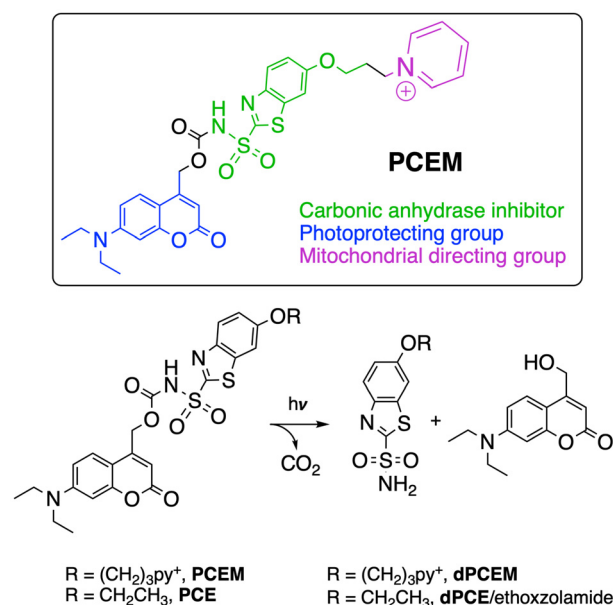


Fig. 1 Design and structure of **PCEM**, a photocaged probe for targeting mitochondrial carbonic anhydrase V isoforms and light-mediated deprotection of **PCEM** and control molecule **PCE**. Note that the structure of **dPCE** is the same as known CA inhibitor ethoxzolamide.

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(coumarin) at one end and linked to a mitochondrial-directing group (pyridinium) at the other end. Aryl sulfonamides are well-established and highly selective inhibitor motifs known to coordinate to the Zn^{2+} ion of CA, thereby obstructing substrate access and inhibiting enzyme activity.^{20,35} The photoprotecting group coumarin exhibits a high quantum yield and a visible light activation wavelength, enhancing its biocompatibility, and can be visualized by fluorescence microscopy.^{36–38} The pyridinium moiety is a delocalized lipophilic cation that targets mitochondria due to its ability to penetrate membranes and its attraction to the negative potential of the mitochondrial matrix.^{39,40} In our approach, the mitochondrial-directing group will shuttle the molecule into the mitochondria as the protecting group simultaneously prevents it from binding to extracellular and cytoplasmic CAs. Once the molecule is within the mitochondria, photoirradiation is used to remove the protecting group, thus exposing the sulfonamide and allowing it to interact with and inhibit mitochondrial CA-V isoforms. To date, this is the first reported photoactivable inhibitor for targeting mitochondrial CA-V isoforms.⁴¹ By providing spatial and temporal control over inhibitor activation, this design enables the selective inhibition of a specific CA isoform within its natural cellular environment.

To optimize our photocaged CA-V inhibitor, we built a homology model for human CA-VA using murine CA-VA (PDB 1URT)⁴² due to their 79% sequence similarity (Fig. S1†). While surface residues showed divergence, active site residues were highly conserved. The active site contains a number of flexible loops; however, the model generated was suitable for docking studies to refine probe design. Using our hCA-VA model, we conducted docking studies to evaluate probe interactions. We first tested triphenyl phosphonium (R-PPh_3^+), a common mitochondrial-targeting group,⁴³ but observed no significant binding regardless of linker length, likely due to steric clashes at the active site entrance. To address this, we replaced R-PPh_3^+ with pyridinium (R-Py^+), which is less bulky and hydrophobic in nature. A three-carbon linker was identified as optimal, balancing Zn^{2+} binding with solubility in aqueous solutions. The docked structure of this final probe design (Fig. 2) shows a coordination bond between the active site Zn^{2+} and the deprotonated sulfonamide moiety, as well as hydrogen bonding interactions with Thr235 and Gln128 in the active site cavity.

In addition to our final photocaged inhibitor **PCEM**, we also synthesized a control molecule, **PCE**, lacking the mitochondrial-targeting group. The full syntheses of both **PCEM** and **PCE** are detailed in Scheme S1† and the deprotection reaction is shown in Fig. 1. The final **PCE** and **PCEM** structures were obtained following coupling of ethoxzolamide or the pyridinium-containing derivative **9** with coumarin **3** in the presence of 4-nitrophenyl chloroformate and DMAP. Irradiation with 360–400 nm light should result in CO_2 release and regeneration of coumarin **3** and the corresponding sulfonamides **dpPCE** and **dpPCEM**. We note that the structures of **dpPCE** and ethoxzolamide are identical, however we will refer to the photodeprotection product of **PCE** as **dpPCE** in characterization

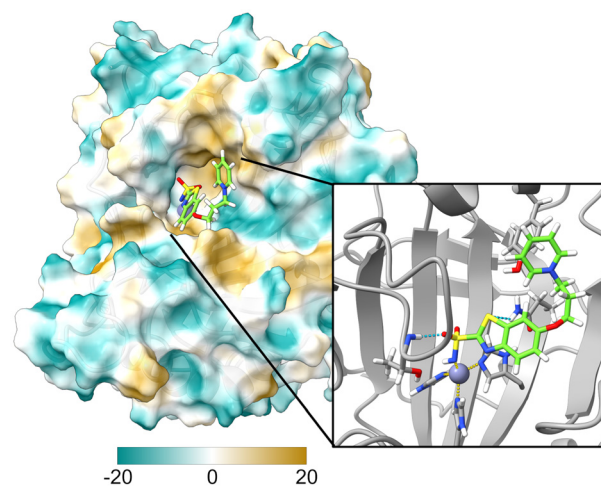


Fig. 2 Docked structure of final **PCEM** probe in hCA-VA homology model. Inset: enlarged view to highlight active site metal coordination (dashed yellow) and hydrogen bonding (dashed blue) interactions.

studies to distinguish samples treated with **PCE** and subsequently deprotected from those directly treated with ethoxzolamide.

The photophysical properties of **PCEM** and **PCE** were assessed in aqueous buffer solution using absorbance and fluorescence spectroscopies and compounds were photoactivated using a Xenon lamp equipped with 365 nm or 410 nm bandpass filters (Fig. S2–S6†). In the absorbance spectrum of **PCE**, a pronounced absorption band at 400 nm was observed, which decreased upon irradiation with either 365 nm or 410 nm light. A similar trend was observed in the fluorescence emission, with the peak at $\lambda_{\text{em}} = 500$ nm ($\lambda_{\text{ex}} = 410$ nm) diminishing slightly following photoirradiation with 365 nm and 410 nm light. We opted to proceed with 410 nm light for further investigations due to higher biocompatibility and slightly faster deprotection kinetics (Fig. S6†). Photodeprotection of **PCE** and **PCEM** and characterization of their resultant products were achieved through LC/MS analysis (Fig. S7–S10†). Conversion of both molecules to coumarin and **dpPCE** or **dpPCEM** was observed without the formation of undesired byproducts.

We used a NPA (4-nitrophenyl acetate) assay⁴⁴ to assess the inhibitory properties of our probes (**PCEM** and **PCE**) and their photo-deprotected analogs (**dpPCEM** and **dpPCE**) after irradiation with 410 nm light. Fig. S11† depicts the increase of absorbance at 400 nm over time when bCA-II was subjected to 1 equivalent of **PCE** and **PCEM** in the absence of light. This change is similar to the control where protein was not treated with any inhibitor, showing the designed probes exhibit no inhibitory properties towards bCA-II. When the probes are irradiated with 410 nm light, to generate **dpPCE** and **dpPCEM** before incubation with bCA-II and NPA, the change in absorbance with time decreases substantially. This trend is similar to the case when bCA-II was treated with ethoxzolamide, proving the inhibitory properties of the probes can be released upon



photoirradiation. For quantitative comparisons between the protected and deprotected forms of the probe, IC_{50} values were assessed (Fig. 3) and compared to ethoxzolamide (IC_{50} = 0.1 μ M). Prior to deprotections, we observed an IC_{50} of 13.6 μ M for **PCE** and much higher for **PCEM** (data could not be fitted to an IC_{50} value). After deprotection, the IC_{50} values decreased to 1.1 μ M for **dPCE** and 2.9 μ M for **dPCEM**, respectively, suggesting that both **dPCE** and **dPCEM** are effective inhibitors for CA, and the presence of the mitochondrial-targeting R-Py⁺ moiety in **dPCEM** does not prevent CA inhibition.

After demonstrating the distinct interaction patterns of **PCE** and **PCEM** with isolated bCA-II protein in both protected and deprotected states, we proceeded to investigate the ability of **PCEM** to target CA-V in isolated mitochondria and in whole cells. Our goals were to demonstrate that our probes are capable of inhibiting the mitochondrial isoforms of CA and show that our combined targeting and photocaging strategy promotes localization of our probes to mitochondria and prevents interaction of our probes with non-mitochondrial CAs.

CAs are essential enzymes that catalyze the conversion of carbon dioxide and water into bicarbonate and protons, playing pivotal roles in pH regulation and CO₂ transport. These processes influence metabolic pathways, including citrulline production in mitochondria. Given that much more is understood about the biology of CA-VA, we focused our studies on this mitochondrial isoform.⁴ It has been reported that inhibition of mitochondrial CA-VA leads to a significant reduction in citrulline levels, likely due to disruption of bicarbonate balance, which affects both the urea cycle and arginine metabolism.⁴⁵ Here, we quantified the total citrulline content in cells using a colorimetric assay in which an indicator reacts with citrulline to form a chromophore and absorbance is measured at 550 nm.⁴⁶ As HepG2 cells express high levels of CA-VA in their mitochondria,¹² testing the ability of our probes to inhibit citrulline production in isolated mitochondria provides a means of assessing the ability of **PCEM** and **PCE** to inhibit this CA isoform. We expect both probes to reduce citrulline production if they are able to effectively bind to the CA-VA active site. We performed a citrulline assay in isolated

mitochondria from HepG2 cells following treatment of cells with 50 μ M **PCE**, **PCEM**, or DMSO (control), alongside 50 μ M ornithine to stimulate citrulline synthesis. At these concentrations, protected and deprotected probes do not show cytotoxicity over a 24 h time period (Fig. S12†). Following treatment with either **PCE** or **PCEM**, cells were irradiated with 410 nm light and then incubated overnight. As shown in Fig. 4A, samples exposed to **dPCE** and **dPCEM** exhibited significantly lower citrulline levels compared to the DMSO control, confirming effective CA inhibition. As expected, no significant difference in citrulline production was observed between **dPCE** and **dPCEM** treatments, which indicates both probes can inhibit CA-VA.

We then studied if addition of the mitochondrial targeting Py⁺ group is effective in directing the probe to mitochondria in whole live cells. For these studies, we focused on HeLa cells, as their flattened morphology in cell culture is more amenable to colocalization studies compared to HepG2 cells, which tend to form clumps of rounded cells.⁴⁷ To confirm localization of **PCEM**, we conducted confocal imaging following the treatment of HeLa cells with MitoTracker Red⁴⁸ and either **PCE** or **PCEM**. As shown in Fig. 4B, **PCE** shows a generalized cytosolic localization, with poor co-localization with MitoTracker (Pearson's R = 0.2, Fig. S13†). Conversely, cells incubated with **PCEM** display more localized staining and higher co-localization with MitoTracker (Pearson's R = 0.4). This correlation difference was statistically significant and supports improved mitochondrial localization of **PCEM** compared to **PCE**.

To demonstrate the differential inhibition properties of **PCEM** and **PCE** in whole cells, we examined how treatment with these probes affects the ability of cytosolic CAs (CA-II) to respond to exogenous CO₂ perturbation. Following CO₂ exposure, extracellular CO₂ is expected to diffuse through the cell membrane, raising its cytoplasmic concentration, where it is converted to carbonic acid by cytoplasmic CAs, resulting in a decrease in cytosolic pH.⁴⁹ Inhibition of cytoplasmic CA reduces the rate of intracellular pH change resulting from the introduction of CO₂.^{50–52} pH changes were tracked over time in the presence of our probes to evaluate real-time inhibition activity. If our mitochondrial targeting strategy is effective, treatment with **PCEM** should not affect the cytosolic pH response.

HeLa cells were treated with pHrodo dye and incubated with **PCE**, **dPCE**, **PCEM**, **dPCEM**, DMSO, and ethoxzolamide. CO₂-saturated water was then introduced into the cell suspension, and fluorescence intensity was monitored over time using flow cytometry, with the changes in pHrodo fluorescence representing variations in intracellular pH. As illustrated in Fig. 4C, CO₂ introduction in the presence of **PCE** and **PCEM** led to a decrease in intracellular pH, similar to the DMSO control. However, after deprotection, **dPCEM** showed no significant change compared to protected **PCEM**, whereas for **dPCE** pH decreased more gradually, similar to ethoxzolamide. This is quantified in Fig. 4D, in which the slopes of the pH decrease are compared. This confirms that introduction of the Py⁺ group effectively localizes **PCEM** and **dPCEM** to the mito-

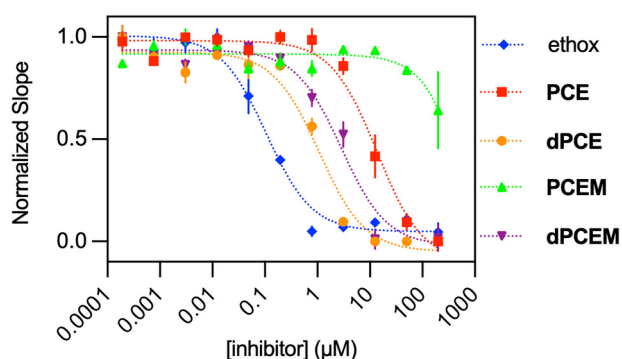


Fig. 3 IC_{50} data for inhibition of NPA hydrolysis by bCA-II in the presence of **PCEM**, **PCE**, **dPCEM**, and **dPCE**. Data for **PCEM** could not be fitted to an IC_{50} value.



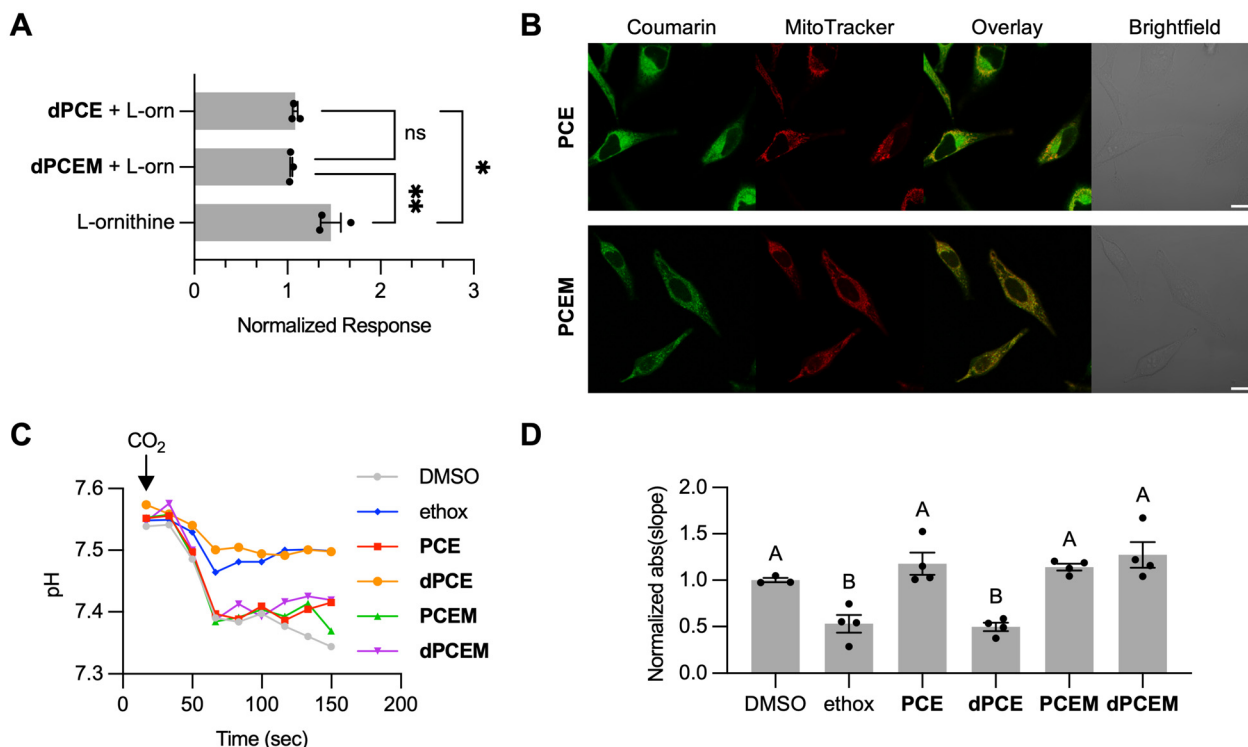


Fig. 4 Cellular studies with PCEM and PCE. (A) Inhibition of citrulline production by dPCE and dPCEM in presence of L-ornithine (B) Confocal images showing localization of PCE and PCEM (green, 1 μ M, $\lambda_{ex/em}$: 405/450–530 nm) and MitoTracker Red (red, 0.1 μ M, $\lambda_{ex/em}$: 560/580–700 nm) in HeLa cells. Scale bar = 20 μ m (C) Change in cytosolic pH over time following addition of CO₂ for cells incubated with DMSO, ethoxzolamide (ethox), (d)PCE or (d)PCEM. (D) Rate of cytosolic pH change following addition of CO₂. The average absolute value of the slope for pH between 33 and 66 s in each condition is shown ($n = 4$). A one-way ANOVA was performed to assess the difference in average slope between all treatments. Bars labeled 'A' and 'B' show groups that are statistically different ($p < 0.05$). All studies were performed in FluoroBrite DMEM at room temperature.

chondria, and thus does not inhibit cytosolic pH change, whereas the untargeted dPCE has a broad CA inhibitory effect, including inhibition of CAs in the cytosol.

Conclusions

We have demonstrated that by combining photocaging and intracellular targeting strategies, a carbonic anhydrase active site inhibitor can be effectively targeted to an intracellular compartment and does not globally inhibit the activity of carbonic anhydrases in cells. Additionally, optimizing probe design to incorporate a sterically compact targeting group allows the photocaged inhibitor to still interact with the protein after activation with light. Next generation designs could incorporate the targeting moiety onto the photoprotecting group, which could improve interactions between the target protein and the deprotected inhibitor. Due to the ease of modifying metal-binding inhibitors with photocaging groups and the abundance of subcellular targeting strategies, we anticipate that this approach will be useful to target other metalloenzymes within many subcellular spaces including mitochondria, the endoplasmic reticulum, the nucleus, and the extracellular membrane.

Author contributions

N. M.: conceptualization, investigation, formal analysis, methodology, validation, visualization, writing – original draft. K. A.: conceptualization, investigation, formal analysis, methodology, validation, visualization, writing – original draft. S. P.: investigation, formal analysis, methodology, validation, visualization, writing – original draft. D. H.: investigation, formal analysis, validation, visualization, writing – review and editing. E. Y.: investigation, methodology, writing – review and editing. D. K.: investigation, methodology, writing – review and editing. E. Q.: conceptualization, formal analysis, visualization, writing – original draft, project administration, resources, funding acquisition, supervision.

Conflicts of interest

There are no conflicts to declare.

Data availability

Data for this article are available at the Texas Data Repository at <https://dataverse.tdl.org/>.



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