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## Role of cyclophilin D-dependent mitochondrial permeability transition in glutamate-induced calcium deregulation and excitotoxic neuronal death

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### Abstract

In the present study we tested the hypothesis that the cyclophilin D-dependent (CyD) mitochondrial permeability transition (CyD-mPT) plays an important role in glutamate-triggered delayed calcium deregulation (DCD) and excitotoxic neuronal death. We used cultured cortical neurons from wild-type C57BL/6 and cyclophilin D knockout mice (*Ppif*<sup>-/-</sup>). Induction of the mPT was identified by following the rapid secondary acidification of mitochondrial matrices monitored with mitochondrially targeted pH-sensitive yellow fluorescent protein. Suppression of the CyD-mPT due to genetic CyD ablation deferred DCD and mitochondrial depolarization, and increased the survival rate after exposure of neurons to 10 $\mu$ M glutamate, but not to 100 $\mu$ M glutamate. Ca<sup>2+</sup> influx into *Ppif*<sup>-/-</sup> neurons was not diminished in comparison with WT neurons judging by <sup>45</sup>Ca accumulation. In both types of neurons, 100 $\mu$ M glutamate produced greater Ca<sup>2+</sup> influx than 10 $\mu$ M glutamate. We hypothesize that greater Ca<sup>2+</sup> influx produced by higher glutamate rapidly triggered the CyD-independent mPT in both WT and *Ppif*<sup>-/-</sup> neurons equalizing their responses to supra-physiologic excitotoxic insults. In neurons exposed to moderate but pathophysiologically-relevant glutamate concentrations, an induction of the CyD-mPT appears to play an important role in mitochondrial injury contributing to DCD and cell death.

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In various neurodegenerations, in traumatic brain injury and stroke, prolonged exposure of neurons to glutamate causes massive Ca<sup>2+</sup> influx into the cytosol (Choi, 1988; Manev et al., 1989; Tymianski et al., 1993b). This produces a rapid jump in the cytosolic Ca<sup>2+</sup> concentration ([Ca<sup>2+</sup>]<sub>c</sub>) followed by its transient decrease to a lower level (Nicholls and Budd, 2000). After some delay, this decrease in [Ca<sup>2+</sup>]<sub>c</sub> is followed by a secondary sustained elevation of [Ca<sup>2+</sup>]<sub>c</sub> or “delayed calcium deregulation” (DCD) (Tymianski et al., 1993a; Nicholls and Budd, 1998). Elevated [Ca<sup>2+</sup>]<sub>c</sub> activates Ca<sup>2+</sup>-dependent degradation enzymes and represents a serious danger to neurons by promoting neuronal death (Wu et al., 2004; Bano et al., 2005; Xu et al., 2007). Therefore, DCD is considered not only a hallmark of but also a potential mechanism leading to glutamate excitotoxicity (Manev et al., 1989; Randall and Thayer,

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1992;Tymianski et al., 1993b). Despite obvious significance and extensive studies, the precise mechanisms leading to DCD still are not completely understood.

Mitochondria accumulate cytosolic  $\text{Ca}^{2+}$  and thus contribute to the clearance of elevated  $[\text{Ca}^{2+}]_c$  (Kiedrowski and Costa, 1995;Wang and Thayer, 1996;White and Reynolds, 1997). However,  $\text{Ca}^{2+}$  uptake can lead to an induction of the mitochondrial permeability transition (mPT) pore accompanied by depolarization of organelles, and therefore, inhibition of further  $\text{Ca}^{2+}$  accumulation (Bernardi, 1999). The molecular composition of the mPT pore is not yet clear, but it is well established that mitochondrial cyclophilin D (CyD) is a regulatory component of the pore (Baines et al., 2005;Basso et al., 2005;Schinzel et al., 2005). It is hypothesized that following excessive  $\text{Ca}^{2+}$  influx into mitochondria, CyD facilitates activation/assembly of the mPT pore (Rasola and Bernardi, 2007). In addition, CyD binds cyclosporin A (CsA), an inhibitor of the CyD-dependent mPT (CyD-mPT) (Crompton et al., 1988). However, under greater  $\text{Ca}^{2+}$  loading, pore activation/assembly may occur in a CyD-independent manner producing the CyD-independent mPT insensitive to CsA (Brustovetsky and Dubinsky, 2000).

In early studies, CsA and its non-immunosuppressive analog, *N*-methyl-valine-4-cyclosporin A (MetVal<sup>4</sup>-CsA), were found to be protective against DCD and/or neuronal death in experiments with oxygen-glucose deprivation (OGD) and in glutamate-treated neurons (Schinder et al., 1996;Nieminen et al., 1996;White and Reynolds, 1996;Khaspekov et al., 1999;Vergun et al., 1999;Almeida and Bolanos, 2001;Alano et al., 2002). CsA and its non-immunosuppressive derivative NIM811 appeared to be protective against neuronal death in traumatic brain and spinal cord injury emphasizing the key role of the mPT pore in these conditions (Sullivan et al., 1999;Scheff and Sullivan, 1999;Sullivan et al., 2000;Okonkwo et al., 1999;Sullivan et al., 2005;McEwen et al., 2007;Mbye et al., 2008). In addition, in a recent study, NIM811, protected against neuronal death in the model of transient focal cerebral ischemia, suggesting involvement of the CyD-mPT (Korde et al., 2007). These studies linked glutamate-induced DCD and neuronal death following ischemic insult to the CyD-mPT. However, other investigators have failed to confirm protective effects of CsA or MetVal<sup>4</sup>-CsA (Isaev et al., 1996;Castilho et al., 1998;Reynolds, 1999;Chinopoulos et al., 2004;Pivovarova et al., 2004). Recently, homozygous knockout mice lacking CyD (*Ppif*<sup>-/-</sup> mice) were generated, and it was shown that isolated mitochondria from these mice have increased resistance to  $\text{Ca}^{2+}$  (Baines et al., 2005;Forte et al., 2007;Nakagawa et al., 2005). Cultured cortical neurons from *Ppif*<sup>-/-</sup> mice appeared to be more resistant to oxidative stress than cells from wild-type animals (WT) (Forte et al., 2007). In addition, *Ppif*<sup>-/-</sup> mice revealed a reduction in brain infarct size after acute middle cerebral artery occlusion (Schinzel et al., 2005). All these effects were attributed to the suppression of the CyD-mPT.

In the present study, we investigated the role of the CyD-mPT in glutamate-triggered DCD and excitotoxic cell death in cultured cortical neurons derived from *Ppif*<sup>-/-</sup> mice and their genetic background C57BL/6 mice. Our experiments revealed that cultured neurons from *Ppif*<sup>-/-</sup> mice were more resistant to glutamate-triggered DCD and cell death than neurons from WT animals. This protection was limited to a moderate but pathophysiologically-relevant glutamate concentration. At higher concentrations, glutamate produced similar DCD and cell death in both *Ppif*<sup>-/-</sup> and WT neurons.

## Experimental Procedures

### Materials

Carbonyl cyanide p-(trifluoromethoxy)phenylhydrazone (FCCP), cyclosporin A (CsA), ADP, oligomycin, glutamate, and glycine were purchased from Sigma (St. Louis, MO). Fura-2FF

AM was bought from Teflabs (Austin, TX) and Rhodamine-123 (FluoroPure™ grade) was purchased from Invitrogen (Carlsbad, CA).

### Isolation and purification of brain mitochondria

Mitochondria from the brains of three cyclophilin D-knockout *Ppif*<sup>-/-</sup> mice (from Dr. Jeffery Molkenstin, University of Cincinnati) and from the brains of three wild-type C57BL/6 mice (Harlan, Indianapolis, IN) were isolated in mannitol-sucrose medium, according to an IACUC approved protocol, and purified on a discontinuous Percoll gradient (Brustovetsky et al., 2002). Mitochondrial protein was determined by the Bradford method (Bradford, 1976) using BSA as a standard. Mitochondria isolated from C57/BL6 and *Ppif*<sup>-/-</sup> mice had a respiratory control index of  $8.27 \pm 0.47$ , N=4, and  $8.53 \pm 0.31$ , N=3, respectively. Respiratory control index is the ratio of the respiratory rate stimulated by addition of 200  $\mu$ M ADP (State 3) to the respiratory rate after inhibition of ADP phosphorylation with 1  $\mu$ M oligomycin (State 4). Mitochondrial respiration was measured using a Clark-type oxygen electrode at 37°C in the standard incubation medium containing 125 mM KCl, 0.5 mM MgCl<sub>2</sub>, 3 mM KH<sub>2</sub>PO<sub>4</sub>, 3 mM glutamate plus 1 mM malate, 10  $\mu$ M EGTA, 0.1% BSA (free from fatty acids), 10 mM HEPES, pH 7.4.

### Slow Ca<sup>2+</sup> infusion

The experiments with slow Ca<sup>2+</sup> infusion were performed as described previously (Chalmers and Nicholls, 2003; Shalbuyeva et al., 2007) with some modifications. Briefly, mitochondria were incubated in the standard incubation medium supplemented with 200  $\mu$ M ADP and 1  $\mu$ M oligomycin in a 0.3 ml chamber at 37°C under continuous stirring. Mitochondrial Ca<sup>2+</sup> accumulation was followed by monitoring disappearance of Ca<sup>2+</sup> from the incubation medium with a miniature Ca<sup>2+</sup> selective electrode. A solution of CaCl<sub>2</sub> was infused into the chamber at a constant rate of 330 nmol CaCl<sub>2</sub>/mg protein  $\times$  min using a KDS 100 pump (KD Scientific, Holliston, MA) equipped with a Hamilton microsyringe. The Ca<sup>2+</sup> uptake capacity was estimated as the amount of accumulated Ca<sup>2+</sup> ( $\mu$ mol per mg of mitochondrial protein). The Ca<sup>2+</sup> uptake capacity was assessed by linear fitting of the early fragment of the experimental trace and the final linear fragment of the trace and then finding the intersection point of these linear graphs (Fig. 1c).

### Western blot analysis of CyD

Isolated brain mitochondria pretreated with Protease Inhibitor Cocktail (Roche, Indianapolis, IN) were solubilized by incubation in a NuPAGE LDS sample buffer (Invitrogen, Carlsbad, CA) supplemented with a reducing agent at 70°C for 15 minutes. 4-12% of the Bis-Tris MOPS gels (Invitrogen) were used for electrophoresis (15  $\mu$ g protein/lane). After electrophoresis, the proteins were transferred to a Hybond™-ECL™ nitrocellulose membrane (Amersham Biosciences, Piscataway, NJ). Blots were incubated for an hour at room temperature with mouse monoclonal anti-cyclophilin D antibody (1:1000 dilution) (EMD Chemicals, San Diego, CA) or goat polyclonal anti-VDAC1 antibody (1:500) (Santa Cruz Biotechnology, Santa Cruz, CA). VDAC1 was used as a control for loading. Blots were developed using goat anti-mouse or donkey anti-goat IgG (1:20000) coupled with horseradish peroxidase (Jackson ImmunoResearch Labs, West Grove, PA) and Supersignal West Pico chemiluminescent reagents (Pierce, Rockford, IL). The molecular weight marker SeeBlue® Plus 2 Standard (5  $\mu$ l) (Invitrogen, Carlsbad, CA) was used to determine the molecular weights of the bands.

### Genotyping

All *Ppif*<sup>-/-</sup> mice were genotyped to ensure that they were homozygous. After preparing the tail samples, the genomic DNA was re-suspended and hydrated in 75  $\mu$ l of Tris-EDTA buffer, pH 8.0. Then, the samples were diluted 1:20 and 1  $\mu$ l of this diluted solution was used for the PCR

(50 $\mu$ l reaction volume). We used three primers in a single reaction: Exon3-F: CTC TTC TGG GCA AGA ATT GC; Neo-F: GGC TGC TAA AGC GCA TGC TCC; and Exon4-R: ATT GTG GTT GGT GAA GTC GCC. The reaction conditions were 95°C for 3 min, then 35 cycles at 95°C for 0.5 min, 56°C for 0.5 min, 72°C for 1 min, and then 72°C for 10 min. The wild-type allele was amplified as a band ~850bp, and the null allele was amplified as a band ~600bp.

### Cell cultures

Primary cultures of cortical neurons were prepared from postnatal day 1 mouse pups according to IACUC approved protocols and procedures previously published for the hippocampus (Dubinsky, 1993), but without preplated glia and the use of antibiotics. For fluorescence measurements, neurons were plated on glass bottomed Petri dishes (Dubinsky, 1993). For all platings, 35  $\mu$ g/ml uridine plus 15  $\mu$ g/ml 5-fluoro-2'-deoxyuridine were added 24 hours after plating to inhibit proliferation of non-neuronal cells. Cultures were maintained in a 5% CO<sub>2</sub> atmosphere at 37°C in Eagle's MEM supplemented with 10% NuSerum (BD Bioscience, Bedford, MA) and 27 mM glucose. Experiments were performed on neurons at 12-14 days *in vitro*.

### Cyclophilin D immunocytochemistry and visualization of mitochondria with MitoTracker Red

To co-localize CyD immunostaining with mitochondria, neurons from C57BL/6 and *Ppif*<sup>-/-</sup> mice, respectively, were stained with MitoTracker Red (Molecular Probes, Eugene, OR). Prior to fixation for immunocytochemistry, neurons were incubated with 0.3 $\mu$ M MitoTracker Red in the growth medium for 15 minutes at 37°C. Then, cultured neurons were fixed with 0.05% glutaraldehyde for 20 minutes, followed by incubation with 4% paraformaldehyde for 15 minutes, and then washed with PBS. Next, cells were incubated with 0.1% glycine-PBS for 30 minutes and incubated with a blocking solution containing 2.5% IgG- and protease-free BSA (Jackson ImmunoResearch Laboratories, West Grove, PA), 2.5% goat serum, and 0.1% Triton X-100 in PBS for an hour at room temperature. Cells were incubated overnight at 4°C with the primary mouse monoclonal anti-cyclophilin D antibody (1:500 dilution) (EMD Chemicals, San Diego, CA). Then, cells were incubated with a secondary donkey anti-mouse antibody conjugated with AlexaFluor 488 (1:1000 dilution) (Invitrogen, Carlsbad, CA) for an hour at room temperature. Bright field and fluorescence images were acquired using a Nikon Eclipse TE2000-U inverted microscope equipped with a Nikon CFI Plan Apo 100 $\times$  1.4 NA objective and Photometrics cooled CCD camera CoolSNAP<sub>HQ</sub> (Roper Scientific, Tucson, AZ) controlled by MetaMorph 6.3 software (Molecular Devices, Downingtown, PA).

### Calcium imaging and monitoring of mitochondrial membrane potential ( $\Delta\psi$ ) in cultured neurons

Cortical neurons were loaded at 37°C simultaneously with 2.6 $\mu$ M Fura-2FF-AM ( $K_d=5.5\mu$ M,  $\lambda_{ex}$  340, 380 nm/ $\lambda_{em}$  512 nm, Molecular Probes, Eugene, OR), to follow changes in cytosolic Ca<sup>2+</sup>, and 1.7 $\mu$ M Rhodamine-123 (Rh123) ( $\lambda_{ex}$  507 nm/ $\lambda_{em}$  529 nm, Molecular Probes, Eugene, OR) to monitor changes in mitochondrial membrane potential ( $\Delta\psi$ ) in the standard bath solution containing 139 mM NaCl, 3 mM KCl, 0.8 mM MgCl<sub>2</sub>, 1.8 mM CaCl<sub>2</sub>, 10 mM NaHEPES, pH 7.4, 5 mM glucose, and 65 mM sucrose. The ion composition of the bath solution is similar or close to those used previously in studies of DCD and excitotoxicity (Wang and Thayer, 1996; White and Reynolds, 1996; Dubinsky et al., 1995; Kushnareva et al., 2005). Sucrose was used to maintain osmolarity similar to that in the growth medium (340 mosm). Osmolarity of the bath solution was measured with an osmometer, Osmette II<sup>TM</sup> (Precision Systems Inc., Natick, MA). Fluorescence imaging was performed with an inverted microscope, Nikon Eclipse TE2000-U, using a Nikon CFI Plan Fluor 20 $\times$  0.45 NA objective and a back-thinned EM-CCD camera Hamamatsu C9100-12 (Hamamatsu Photonic Systems, Bridgewater, NJ) controlled by Simple PCI software 6.1 (Compix Inc., Sewickley, PA). The

excitation light was delivered by a Lambda-LS system (Sutter Instruments, Novato, CA). The excitation light at 480 nm was attenuated by quartz neutral density filters to 10%. The excitation filters ( $340\pm 5$ ,  $380\pm 7$ , and  $480\pm 20$ ) were controlled by a Lambda 10-2 optical filter changer (Sutter Instruments, Novato, CA). Fluorescence was recorded through a 505 nm dichroic mirror at  $535\pm 25$  nm. To minimize photobleaching and phototoxicity, the images were taken every 15 seconds during the time-course of the experiment using the minimal exposure time that provided acceptable image quality. The changes in  $[Ca^{2+}]_c$  were monitored by following a ratio of  $F_{340}/F_{380}$ , calculated after subtracting the background from both channels. The changes in the  $\Delta\psi$  were monitored by following changes in the fluorescence of Rh123 expressed as  $F/F_0$ . The Rh123 fluorescence traces were also constructed after subtracting the background. The contribution of plasma membrane depolarization to the Rh123 signal appeared to be negligible as determined by applying  $Ca^{2+}$ -free bath solution containing 50 mM KCl and a correspondingly decreased concentration of NaCl to Rh123-loaded neurons (not shown). After 3 minutes of fluorescence recording in the standard bath solution, various concentrations of glutamate (10 or 100  $\mu$ M) plus 10  $\mu$ M glycine were applied to the neurons. At the end of the experiment, the bath solution with glutamate and  $Ca^{2+}$  was replaced by a glutamate- and  $Ca^{2+}$ -free solution, and then 1  $\mu$ M FCCP was applied to the neurons to depolarize neuronal mitochondria and to release  $Ca^{2+}$  accumulated in mitochondria.

### Transfection of primary cortical neurons

To measure matrix pH ( $pH_m$ ) in mitochondria within live cells, hippocampal neurons were transfected in suspension during plating using an electroporator BTX 630 ECM (Harvard Apparatus, Holliston, MA) with a plasmid encoding mitochondrially targeted eYFP (enhanced Yellow Fluorescent Protein, generously provided by Dr. Roger Tsien, University of California, San Diego) as described previously (Shalbuyeva et al., 2006). Neurons were taken into the experiment 12-14 days after transfection.

### Confocal microscopy

Laser-based spinning-disk confocal microscopy was used to visualize the mitochondrial network in neurons expressing mito-eYFP. The imaging was performed with an inverted microscope, Nikon Eclipse TE2000-U, equipped with spinning-disk confocal unit Yokogawa CSU-10 (Yokogawa Electric Corp. Tokyo, Japan) using a Nikon CFI Plan Apo 100 $\times$  1.3 NA objective and a back-thinned EM-CCD camera Andor iXon+ DU-897 (Andor, Morrisville, NC) controlled by Andor iQ (Andor, Morrisville, NC). A 2 $\times$  extender was placed in front of the camera to increase spatial resolution. The serial images (z-stack) were acquired with a z-step 0.1  $\mu$ m using a piezo-electric positioning device PIFOC P-721 (Physik Instrumente, Karlsruhe, Germany). Image processing consisted of a blind 3D deconvolution using AutoQuant X 2.1.1 (MediaCybernetics, Bethesda, MD) and a 3D rendering using Imaris 5.7.2 (Bitplane, St. Paul, MN).

### Matrix pH measurements

Matrix pH ( $pH_m$ ) measurements and pH calibration were performed as described previously (Bolshakov et al., 2008). In addition, neurons expressing mito-eYFP were loaded with Fura-2FF to follow  $pH_m$  and changes in cytosolic  $Ca^{2+}$  concentration simultaneously. In these experiments we used a Nikon Eclipse TE2000-U inverted microscope equipped with a Nikon CFI SuperFluor 40 $\times$  1.3 NA objective and a Photometrics cooled CCD camera CoolSNAP<sub>HQ</sub> (Roper Scientific, Tucson, AZ) controlled by MetaMorph 6.3 software (Molecular Devices, Downingtown, PA).

## <sup>45</sup>Ca accumulation

Experiments were performed as described previously (Hartley et al., 1993) with some modifications. Cortical neurons from WT or *Ppif*<sup>-/-</sup> mice were plated on a 12-well plate. The neurons were washed once with standard bath solution to remove growth medium. Then, neurons were incubated with a solution containing 10 or 100 μM glutamate (plus 10 μM glycine) and supplemented with 5 μCi/ml <sup>45</sup>Ca. The neurons were then washed three times in an ice-cold, Ca<sup>2+</sup>-free bath solution. Neurons were lysed with 0.5% SDS. An aliquot was added to the scintillation liquid Ecolite (MP Biomedicals, Santa Ana, CA) and counted in a Tri-Carb 2100TR liquid scintillation analyzer (Packard Instrument Co., Meriden, CT). Each condition was carried out three times.

## Glutamate toxicity

After culturing mouse cortical neurons for 12-14 DIV, the medium was replaced with Eagle's MEM without serum supplemented with 27 mM glucose, 15 mM sucrose, 100 μM glutamine, and 10 μM glycine (Brustovetsky et al., 2004). The neurons were then exposed to various concentrations of glutamate (3-300 μM) for 10 minutes. After that, glutamate was removed, and the cells were rinsed with Eagle's Balanced Salt Solution (EBSS) supplemented with 27 mM glucose and 15 mM sucrose, and left in the thermostat at 37°C for the next 24 hours. After 24 hours, cell death was assessed by nuclear staining with 4.5 μM propidium iodide (PI) (Pivovarov et al., 2004). Nuclei staining with PI is associated with the loss of barrier properties of the plasma membrane and is considered an indication of necrosis (Orrenius et al., 2003). In addition, neuronal death was quantitatively evaluated with the Trypan Blue exclusion method (Dubinsky and Rothman, 1991). An induction of apoptosis was evaluated with Annexin V staining (Molecular Probes, Eugene, OR). Dying neurons were detected using a Nikon Eclipse TE2000-U inverted microscope equipped with a Nikon CFI SuperFluor 20× 0.75 NA objective and a Photometrics cooled CCD camera CoolSNAP<sub>HQ</sub> (Roper Scientific, Tucson, AZ) controlled by MetaMorph 6.3 software (Molecular Devices, Downingtown, PA). These toxicity experiments were also performed in triplicate on neurons from three separate platings.

## Statistics

Every experiment was performed using at least three separate preparations of isolated mitochondria or three independent, separate neuronal platings. All data represent mean ± SEM of at least 3 separate and independent experiments. Statistical analysis of the experimental results consisted of a one-way ANOVA followed by Bonferroni's *post hoc* test (GraphPad Prism<sup>®</sup> 4.0, GraphPad Software Inc., San Diego, CA). *t*-scores calculated in GraphPad Prism<sup>®</sup> are shown in the legends to the Figures with statistical analysis.

## Results

### Calcium uptake capacity of isolated brain mitochondria from C57BL/6 and *Ppif*<sup>-/-</sup> mice

In our experiments, we used homozygous *Ppif*<sup>-/-</sup> mice lacking CyD (Baines et al., 2005) to test the hypothesis that the CyD-mPT contributes to mitochondrial injury in neurons exposed to glutamate, thus limiting the ability of the organelles to participate in the maintenance of [Ca<sup>2+</sup>]<sub>c</sub>. All mice used in our experiments were genotyped using PCR analysis of genomic DNA. Figure 1a shows the results of genotyping two C57BL/6 mice and two *Ppif*<sup>-/-</sup> mice. C57BL/6 mice are genetic background for *Ppif*<sup>-/-</sup> mice. Therefore, we used isolated mitochondria and cultured neurons derived from C57BL/6 mice as controls for the experiments with mitochondria and neurons derived from *Ppif*<sup>-/-</sup> mice. The wild-type allele amplifies as a band ~850bp and the null allele amplifies as a band ~600bp as shown. Western blotting analysis confirmed the lack of CyD in brain mitochondria isolated from *Ppif*<sup>-/-</sup> mice (Fig. 1b). Consistent with previous reports (Basso et al., 2005; Nakagawa et al., 2005; Schinzel et al., 2005; Forte et

al., 2007; Naga et al., 2007), in the experiments with slow  $\text{CaCl}_2$  infusion, brain mitochondria isolated from *Ppif*<sup>-/-</sup> mice had greater  $\text{Ca}^{2+}$  uptake capacity than mitochondria from WT mice (Fig. 1c). In this study, we used non-synaptic mitochondria, which mostly represented a mixture of mitochondria from glial cells and neuronal somata. In our experiments with cultured neurons, we were focused on perturbations in cytosolic  $\text{Ca}^{2+}$  and mitochondrial membrane potential in the neuronal somata. Therefore, isolated non-synaptic mitochondria were well-suited to our purpose. CsA (1  $\mu\text{M}$ ) significantly increased  $\text{Ca}^{2+}$  uptake capacity of mitochondria from WT mice (Fig. 1d) implicating the CyD-mPT as a mechanism that limits the ability of mitochondria to accumulate  $\text{Ca}^{2+}$ . On the other hand, CsA was without effect in mitochondria from *Ppif*<sup>-/-</sup> mice where CyD, the target for CsA, was genetically ablated (Fig. 1e). Figure 1f shows the summary of data obtained with mitochondria from wild-type and *Ppif*<sup>-/-</sup> mice and incubated with or without 1  $\mu\text{M}$  CsA. Thus, isolated brain mitochondria from *Ppif*<sup>-/-</sup> mice with suppressed CyD-mPT demonstrated higher resistance to  $\text{Ca}^{2+}$  than mitochondria from WT animals. However, the increasing  $\text{Ca}^{2+}$  load eventually damaged mitochondria from both *Ppif*<sup>-/-</sup> and WT mice and precluded further  $\text{Ca}^{2+}$  accumulation probably due to the induction of the CyD-independent mPT.

### Delayed calcium deregulation in cortical neurons from C57BL/6 and *Ppif*<sup>-/-</sup> mice

Next, we addressed the question of whether cultured cortical neurons derived from *Ppif*<sup>-/-</sup> mice have an increased resistance to glutamate-induced DCD in comparison with neurons from WT animals. To confirm the lack of CyD expression in mitochondria of cultured neurons, we performed an immunocytochemistry analysis using an antibody against CyD. To ensure mitochondrial localization of CyD, neurons were co-stained with a mitochondrial marker, MitoTracker Red, and then CyD and MitoTracker Red images were overlaid (Fig. 2). Immunostaining of cultured neurons derived from WT animals revealed an abundance of CyD in mitochondria, while in neurons from *Ppif*<sup>-/-</sup> mice CyD was below the detection limit. To evaluate the role of the CyD-mPT in glutamate-induced DCD, neurons were co-loaded with calcium-sensitive, low-affinity fluorescence dye Fura-2FF and mitochondrial membrane potential-sensitive dye Rhodamine-123 (Rh123). Figure 3 shows Fura-2FF and Rh123 fluorescence traces from the representative experiments with cultured neurons derived from WT (a,c) and *Ppif*<sup>-/-</sup> mice (b,d). The individual traces (thin grey lines) were obtained from different individual neurons and overlapped by average traces (thick black lines for Fura-2FF and for Rh123, mean  $\pm$  SEM). Following a few minutes of incubation in the standard bath solution, neurons were exposed to 10  $\mu\text{M}$  glutamate plus 10  $\mu\text{M}$  glycine for 20 minutes. This concentration of glutamate was chosen based on early data obtained after cerebral ischemia in rats (Benveniste et al., 1984; Van Hemelrijck et al., 2005) and on clinical reports indicating an increase of glutamate concentration in the cerebrospinal fluid of stroke patients up to 7-8  $\mu\text{M}$  (Castillo et al., 1997; Castillo et al., 1996). After 20 minutes of incubation, the glutamate and  $\text{Ca}^{2+}$ -containing bath solution was replaced with a glutamate- and  $\text{Ca}^{2+}$ -free solution to assess the ability of neurons to recover  $[\text{Ca}^{2+}]_c$ . Finally, 1  $\mu\text{M}$  FCCP was applied to neurons to completely depolarize the entire mitochondrial population and release accumulated  $\text{Ca}^{2+}$ .

In these experiments, both WT and *Ppif*<sup>-/-</sup> neurons responded to glutamate with a fast initial jump in  $[\text{Ca}^{2+}]_c$  followed by a decrease to a new, somewhat elevated level (Fig. 3). Soon after that, a secondary slower but greater increase in  $[\text{Ca}^{2+}]_c$  took place in WT neurons indicating the onset of DCD (Fig. 3a). The recovery of  $[\text{Ca}^{2+}]_c$  after the removal of glutamate and  $\text{Ca}^{2+}$  was slow in WT neurons, and the release of  $\text{Ca}^{2+}$  from mitochondria in response to FCCP-induced mitochondrial depolarization was negligible (Fig. 3a). Simultaneous to the increases in  $[\text{Ca}^{2+}]_c$  we observed two waves of mitochondrial depolarization manifested in the increase of Rh123 fluorescence. Interestingly, at the end of the experiments, FCCP increased the Rh123 signal presumably by depolarizing uninjured mitochondria. This hypothesis is supported by the observations that mitochondria are functionally heterogeneous within the cell (Collins et

al., 2002) and that glutamate-treated neurons may contain both damaged and normal, uninjured mitochondria at the same time (Pivovarova et al., 2004). Alternatively, it could be an artifact due to pH change in the cytosol and in the mitochondrial matrix. However, FCCP did not produce significant changes in cytosolic or matrix pH following DCD in cultured neurons exposed to glutamate (Bolshakov et al., 2008), therefore this scenario seems unlikely.

Mitochondria significantly contribute to maintenance of calcium homeostasis in neuronal cells (Herrington et al., 1996; Pivovarova et al., 2004). Mitochondrial damage due to induction of the CyD-mPT might cause depolarization of the organelles precluding further  $\text{Ca}^{2+}$  accumulation by mitochondria. Therefore, we hypothesized that protection of mitochondria from the CyD-mPT by genetic ablation of CyD could protect mitochondria, enhance their ability to accumulate  $\text{Ca}^{2+}$  and thereby increase resistance of neurons to DCD and secondary mitochondrial depolarization. Indeed, in *Ppif*<sup>-/-</sup> neurons lacking CyD, the induction of DCD and secondary mitochondrial depolarization were significantly deferred (Fig. 3b). However, even *Ppif*<sup>-/-</sup> neurons eventually experienced DCD and secondary mitochondrial depolarization. This is consistent with the facts that CyD ablation does not provide an absolute protection against  $\text{Ca}^{2+}$ -induced mitochondrial damage (Fig. 1c) and mPT may occur in a CyD-independent manner (Basso et al., 2005).

In some pathological conditions, the concentration of glutamate in the extracellular milieu might jump up to 100 $\mu\text{M}$  within a few minutes of global ischemia, leading to severe neuronal injury and cell death (Benveniste et al., 1984; Nakayama et al., 2002). In our next experiments, we investigated whether the protective effects of CyD ablation could be extended to a greater glutamate challenge. It appeared that both WT and *Ppif*<sup>-/-</sup> neurons responded similarly to 100 $\mu\text{M}$  glutamate (Fig. 3c,d). Thus, the protective role of CyD ablation was evident only with a moderate glutamate concentration.

In the experiments with calcium and  $\Delta\psi$  imaging, different neurons in the same dish responded to glutamate somewhat differently (Fig. 3). This is a typical behavior of individual cultured neurons (Schinder et al., 1996; Nieminen et al., 1996; White and Reynolds, 1996; Khaspekov et al., 1999; Vergun et al., 1999; Almeida and Bolanos, 2001; Alano et al., 2002; Isaev et al., 1996; Castilho et al., 1998; Reynolds, 1999; Chinopoulos et al., 2004; Pivovarova et al., 2004; Manev et al., 1989; Tymianski et al., 1993b; Thayer and Miller, 1990; Budd and Nicholls, 1996). In addition, in our experiments we observed some variations in responses of neurons from different platings. To provide a statistical analysis of the data, we introduced a parameter: *the time from the beginning of glutamate exposure to the completion of the DCD* ( $t_{DCD}$ ) (Fig. 3a,b). A similar approach was used previously to analyze secondary mitochondrial depolarization in cultured neurons exposed to glutamate (Vergun et al., 2003). Our statistical analysis confirmed that glutamate triggered DCD more rapidly in WT neurons than in *Ppif*<sup>-/-</sup> neurons (Fig. 4). At the same time,  $t_{DCD}$  was similar both in WT and in *Ppif*<sup>-/-</sup> neurons exposed to 100 $\mu\text{M}$  glutamate. There were 8 separate, independent experiments in each group with neurons from five different platings. N indicates total number of cells in each group. Thus, the statistical analysis of data obtained with calcium imaging showed that the protection against DCD evoked by CyD ablation was restricted to a moderate glutamate concentration and was not evident with a greater glutamate challenge.

The lack of protection due to CyD ablation in neurons exposed to high glutamate could be because of a greater  $\text{Ca}^{2+}$  influx and faster mitochondrial damage. In our experiments with isolated mitochondria, increasing  $\text{Ca}^{2+}$  loading ultimately overcame protection conferred by CsA or by CyD ablation (Fig. 1c-f). To assess the kinetics and the amount of  $\text{Ca}^{2+}$  influx into neurons, we measured <sup>45</sup>Ca accumulation in the cells. Every experiment was performed in triplicate with neurons from three different platings. Glutamate at a higher concentration (100 $\mu\text{M}$ ) caused faster <sup>45</sup>Ca influx into neurons than a lower concentration of glutamate



(10 $\mu$ M) (Fig. 5). Therefore, with higher glutamate mitochondrial damage might occur earlier and the protection against DCD in *Ppif*<sup>-/-</sup> neurons could be significantly diminished or completely vanished. In addition, *Ppif*<sup>-/-</sup> neurons accumulated <sup>45</sup>Ca more rapidly in comparison with WT neurons indicating the lack of decrease in Ca<sup>2+</sup> influx into *Ppif*<sup>-/-</sup> neurons. Thus, it seemed very unlikely that the deferment in DCD and mitochondrial depolarization in *Ppif*<sup>-/-</sup> neurons exposed to moderate glutamate was due to attenuation of the Ca<sup>2+</sup> influx into the cells.

Recently, a complex dynamics of mitochondrial matrix acidification in neurons exposed to glutamate was revealed with pH-sensitive enhanced yellow fluorescent protein targeted to mitochondria (mito-eYFP) (Bolshakov et al., 2008). In these experiments, the second phase of matrix acidification following glutamate application reflected an induction of the mPT pore leading to a drastic increase in ion permeability of the inner mitochondrial membrane and pH equilibration between the cytosol (more acidic) and the mitochondrial matrix (initially more alkaline). In the current study, we used this approach in the experiments with WT and *Ppif*<sup>-/-</sup> neurons to demonstrate the relationship between the onset of DCD and the induction of the CyD-mPT. We examined 39-48 individual neurons in each group in 12 separate, independent experiments with cells from six different platings. Figure 6a shows representative bright field (right) and fluorescent (left) images of neurons expressing mito-eYFP. The expression rate was in the range of 10-15%. Figure 6b demonstrates 3D reconstruction of the mitochondrial network in the neuron expressing mito-eYFP. Figures 6c-e illustrate results from the experiments with WT or *Ppif*<sup>-/-</sup> neurons exposed to 10 $\mu$ M glutamate (plus 10 $\mu$ M glycine). Statistical analysis of these experiments is shown in Figure 6f. The onset of secondary matrix acidification (SMac) was quantitatively assessed in all tested WT neurons and in those *Ppif*<sup>-/-</sup> neurons that experienced SMac within the time course of the experiment. The onset of DCD in WT and *Ppif*<sup>-/-</sup> neurons coincided with SMac. (Fig. 6c,d). In the case with WT neurons, the addition of FCCP at the end of the experiment did not produce a significant change in pH<sub>m</sub>. In contrast to WT neurons, in *Ppif*<sup>-/-</sup> neurons SMac and DCD were significantly deferred in 38 (79.2%) or absent in 10 (20.8%) out of 48 neurons examined in 12 separate experiments (Fig. 6d,e). With *Ppif*<sup>-/-</sup> neurons that did not experience SMac and DCD, FCCP produced significant matrix acidification due to equilibration of pH<sub>m</sub> and cytosolic pH accompanied by a strong increase in [Ca<sup>2+</sup>]<sub>c</sub>, suggesting a massive release of accumulated Ca<sup>2+</sup> from mitochondria given that external Ca<sup>2+</sup> was removed. Such a strong FCCP-induced increase in [Ca<sup>2+</sup>]<sub>c</sub> was not observed after DCD in WT neurons. In both WT and *Ppif*<sup>-/-</sup> neurons, SMac was not observed when neurons were exposed to glutamate in Ca<sup>2+</sup>-free bath solution (not shown). When neurons were exposed to 100 $\mu$ M glutamate, the onset of SMac occurred simultaneously in WT and *Ppif*<sup>-/-</sup> neurons (Fig. 6f). Thus, in neurons exposed to moderate glutamate, mitochondrial damage due to induction of the CyD-mPT substantially contributed to DCD. Suppression of the CyD-mPT induction, and hence preservation of mitochondrial Ca<sup>2+</sup> uptake, at least temporarily protected neurons from the collapse of calcium homeostasis leading to postponement of DCD. With a high level of glutamate, mitochondrial damage occurred in a CyD-independent manner presumably due to induction of the CyD-independent mPT.

### Excitotoxic cell death in cortical neurons from C57BL/6 and *Ppif*<sup>-/-</sup> mice

There is a substantial body of evidence suggesting a causal link between sustained elevation of cytosolic Ca<sup>2+</sup> and excitotoxic neuronal death (Tymianski et al., 1993c; Tymianski et al., 1994; Limbrick, Jr. et al., 1995; Brustovetsky et al., 2004). In our next experiments, we addressed the question of whether suppression of the glutamate-triggered, Ca<sup>2+</sup>-dependent CyD-mPT due to genetic CyD ablation leads to an increase in the survival rate of neurons exposed to glutamate. Glutamate-induced cell death was evaluated following propidium iodide (PI) nuclear staining (Pivovarova et al., 2004). This method has been chosen because it detects necrosis, which significantly contributes to neuronal death following prolonged glutamate

exposure (Ankarcrona et al., 1995). However, there is a possibility that in *Ppif*<sup>-/-</sup> neurons necrosis might be substituted for another type of cell death, apoptosis. To examine this possibility, we double stained neurons with PI, a marker of necrosis, and Annexin V, a marker of the early stage of apoptosis. 10 minute exposure of neurons from WT or *Ppif*<sup>-/-</sup> mice to 10 $\mu$ M glutamate (plus 10 $\mu$ M glycine) 24 hours later caused significant cell death in WT neurons but not in *Ppif*<sup>-/-</sup> neurons (Fig. 7). Thus, CyD ablation resulted in remarkable protection against glutamate toxicity. With both WT and *Ppif*<sup>-/-</sup> neurons, Annexin V staining was negligible. As a positive control for Annexin V staining, we used *Ppif*<sup>-/-</sup> neurons treated for 24 hours with 30 $\mu$ M cisplatin (Cis), an agent that induces apoptosis. Treatment of neurons with cisplatin caused strong Annexin V staining, indicating an induction of apoptosis (Fig. 7). Thus, protection of *Ppif*<sup>-/-</sup> neurons from glutamate-triggered necrotic cell death was not accompanied by increased apoptosis.

In addition, neuronal death was quantitatively assessed across the range of glutamate concentrations (3-300 $\mu$ M) by the Trypan Blue exclusion method (Dubinsky and Rothman, 1991). There were 9 independent experiments with neurons from nine different platings. In each experiment, cells were counted three times in randomly chosen fields and the averages we used for statistical analysis. In these experiments, neurons were exposed to various concentrations of glutamate for 10 minutes. At this time, all WT neurons treated with 10 $\mu$ M glutamate experienced DCD and reached the final elevated  $[Ca^{2+}]_c$  plateau, whereas *Ppif*<sup>-/-</sup> neurons still had a relatively low level of  $[Ca^{2+}]_c$  (Fig. 3a,b). On the other hand, after 10 minutes of exposure to 100 $\mu$ M glutamate, both *Ppif*<sup>-/-</sup> and WT neurons had completed DCD (Fig. 3c,d). Accordingly, the glutamate toxicity experiments revealed greater survival of *Ppif*<sup>-/-</sup> neurons than WT neurons in the range of 10 $\mu$ M glutamate, while at higher glutamate concentrations the survival rate of *Ppif*<sup>-/-</sup> and WT neurons appeared to be similar (Fig. 8). Thus, the protective effect of CyD ablation against excitotoxic neuronal death was found to be significant at moderate but pathophysiologically-relevant concentrations of glutamate, while at higher glutamate concentrations the protection was not evident.

## Discussion

In this study we demonstrated for the first time the key role of CyD-mPT in DCD and excitotoxic neuronal death using cultured neurons derived from CyD-knockout *Ppif*<sup>-/-</sup> mice. In contrast to pharmacological inhibitors which may have off-target effects, genetic ablation of CyD is free of such confounding complications, and therefore represents a more advantageous model for studying the role of the CyD-mPT in glutamate excitotoxicity and other pathologies.

The main finding of our study is that genetic ablation of CyD, which leads to suppression of the CyD-mPT, improves handling of calcium homeostasis and preserves mitochondrial membrane potential in cultured cortical neurons exposed to pathophysiologically-relevant glutamate concentration. Most importantly, CyD ablation increases survival rate of neurons exposed to moderate glutamate, and thus unequivocally demonstrates the important role of the CyD-mPT in glutamate excitotoxicity. However, while suppression of the CyD-mPT due to CyD ablation defers the collapse of calcium homeostasis and improves survival rate of neurons, this genetic manipulation fails to completely protect neurons against glutamate, emphasizing the potential multiplicity of mechanisms contributing to neuronal injury.

Early studies suggested an important role for the mPT in the disturbance of calcium homeostasis and neuronal death in the experimental model of traumatic brain injury (Sullivan et al., 1999; Scheff and Sullivan, 1999; Sullivan et al., 2000; Okonkwo et al., 1999; Sullivan et al., 2005; McEwen et al., 2007; Mbye et al., 2008), in oxygen/glucose deprivation (OGD) and in neurons exposed to excitotoxic glutamate (Schinder et al., 1996; Nieminen et al., 1996; White

and Reynolds, 1996;Almeida and Bolanos, 2001;Alano et al., 2002;Khaspekov et al., 1999;Vergun et al., 1999). In these studies, CsA appeared to be neuroprotective, linking DCD and excitotoxic neuronal injury to induction of the mPT. However, in other studies, pharmacological inhibition of CyD appeared to be futile (Isaev et al., 1996;Castilho et al., 1998;Reynolds, 1999;Chinopoulos et al., 2004;Pivovarova et al., 2004). Since then, the lack of CsA protection against glutamate was used as the main argument against mPT involvement in DCD and excitotoxic neuronal death. The contradictory results obtained in early studies with CsA could be explained by the fact that CsA binds to both CyD and cytosolic cyclophilin A, and the latter leads to inhibition of calcineurin, which has numerous targets in the cell (Liu et al., 1991;Yakel, 1997). *N*-methyl-valine-4-cyclosporin A (MetVal<sup>4</sup>-cyclosporin), a CsA derivative that does not inhibit calcineurin (Zenke et al., 1993) but potently suppresses the mPT (Friberg and Wieloch, 2002), has been used in several studies to distinguish between inhibition of calcineurin and suppression of the mPT (Vergun et al., 1999;Alano et al., 2002;Khodorov, 2004;Castilho et al., 1998). Yet, even with MetVal<sup>4</sup>-cyclosporin, the results appeared to be rather controversial. Some investigators reported that MetVal<sup>4</sup>-cyclosporin is neuroprotective (Vergun et al., 1999;Alano et al., 2002;Khodorov, 2004) while others did not observe protection with this pharmacological agent (Castilho et al., 1998). The reason for this discrepancy is not quite clear. A possible explanation lies in the fact that CsA protection against the mPT is highly variable and strongly depends on experimental conditions. It significantly diminishes or completely vanishes following an increase in the magnitude of Ca<sup>2+</sup> loading, mitochondrial depolarization, increased duration of the insult, or interaction of mitochondria with some agents such as free fatty acids (FFA) (Bernardi et al., 1993;Bernardi et al., 1992;Broekemeier and Pfeiffer, 1989;Broekemeier and Pfeiffer, 1995;Brustovetsky and Dubinsky, 2000).

Consistent with this notion, the protection of neurons against glutamate due to the inhibition of the CyD-mPT achieved by genetic CyD ablation depended on the severity of the glutamate insult. This fact was not established, and therefore was not appreciated in early studies with pharmacological inhibitors of CyD. We found in our experiments that excessive Ca<sup>2+</sup> influx into neurons via activated glutamate receptors readily overrode protection imposed by inhibition of the CyD-mPT achieved by genetic CyD ablation. While neurons from *Ppif*<sup>-/-</sup> mice were more resistant to DCD following exposure to 10μM glutamate, with 100μM glutamate the difference between *Ppif*<sup>-/-</sup> and WT neurons became negligible. Importantly, cultured neurons exposed to 100μM glutamate accumulated <sup>45</sup>Ca faster than neurons exposed to 10μM glutamate. It is conceivable that the faster delivery of Ca<sup>2+</sup> to mitochondria and, correspondingly faster Ca<sup>2+</sup> loading into mitochondria, might more rapidly trigger an induction of the mPT. The greater Ca<sup>2+</sup> influx into neurons exposed to higher glutamate could also rapidly induce the CyD-independent mPT, which damages mitochondria due to dramatic permeabilization of the IMM regardless of the level of CyD expression. This might be accountable for the similar responses of WT and *Ppif*<sup>-/-</sup> neurons to high glutamate. The rapid and excessive Ca<sup>2+</sup> influx into neurons could also be one of the reasons why CsA or its derivatives were not effective in the early experiments with high glutamate concentrations (Isaev et al., 1996;Castilho et al., 1998;Reynolds, 1999;Chinopoulos et al., 2004;Pivovarova et al., 2004).

It has been well established in early studies that an induction of the mPT pore in isolated mitochondria can cause a loss of pyridine nucleotides (Vinogradov et al., 1972;Di et al., 2001). This, in turn, can cause an inhibition of mitochondrial respiration that could be restored either by addition of NAD<sup>+</sup> (Fontaine et al., 1998) or by addition of succinate (Brustovetsky et al., 2002;Brustovetsky et al., 2003). However, the inhibition of mitochondrial respiration due to loss of pyridine nucleotides primarily happens if mitochondria are fueled by Complex I-linked substrates exclusively. This seems quite unlikely in *in situ* conditions when in addition to Complex I-linked substrates mitochondria are also fueled by succinate. Nevertheless, the

loss of pyridine nucleotides via the mPT pore might decrease reserve respiratory capacity of mitochondria thus contributing to mitochondrial dysfunction.

In contrast to the experiments with isolated mitochondria, identifying and assessing the role of the CyD-mPT in the experiments with cultured neurons appeared to be a daunting task. In addition to the use of pharmacological inhibitors of the CyD-mPT, some investigators employed alternative approaches in attempt to better understand the role of the CyD-mPT in disturbances of calcium homeostasis and excitotoxicity. Recently, Kushnareva et al. found increased amounts of  $\text{Ca}^{2+}$  accumulated in mitochondria isolated from cultured cortical neurons exposed to excitotoxic glutamate (Kushnareva et al., 2005). This was interpreted as strong evidence against involvement of the mPT in glutamate-triggered DCD, since the opening of the mPT pore should cause a release of previously accumulated  $\text{Ca}^{2+}$  (Bernardi and Petronilli, 1996). However, in another elegant study, Pivovarova et al., using electron probe X-ray microanalysis, observed significant calcium precipitates in swollen mitochondria within glutamate-treated cultured hippocampal neurons (Pivovarova et al., 2004). The swelling of mitochondria was attributed to the induction of the mPT, and it was proposed that, probably due to poor solubility (Chalmers and Nicholls, 2003), calcium precipitates remain in mitochondria even after the opening of the mPT pore (Pivovarova et al., 2004). Recently, this point of view received additional experimental support in experiments with isolated brain mitochondria (Kristian et al., 2007). Thus, an increased amount of  $\text{Ca}^{2+}$  in mitochondria from glutamate-treated neurons does not necessarily indicate a lack of the mPT, and therefore obviously cannot serve as evidence against mPT induction in neurons exposed to excitotoxic glutamate.

The advent of CyD-knockout *Ppif*<sup>-/-</sup> mice, which were used in the present study, permitted us to address the role of the CyD-mPT in DCD, secondary mitochondrial depolarization, and glutamate excitotoxicity without the use of pharmacological agents. The total elimination of CyD in neuronal mitochondria of *Ppif*<sup>-/-</sup> mice ensured the maximal suppression of the CyD-dependent mPT, which cannot be exceeded by any existing or future pharmacological inhibitors of CyD. The experiments performed in this study provided a deeper insight into the mechanisms of glutamate-triggered DCD and excitotoxicity and demonstrated that the use of neurons from *Ppif*<sup>-/-</sup> mice represents a new, valuable approach in examining the role of the CyD-mPT in pathophysiological processes including DCD and glutamate excitotoxicity.

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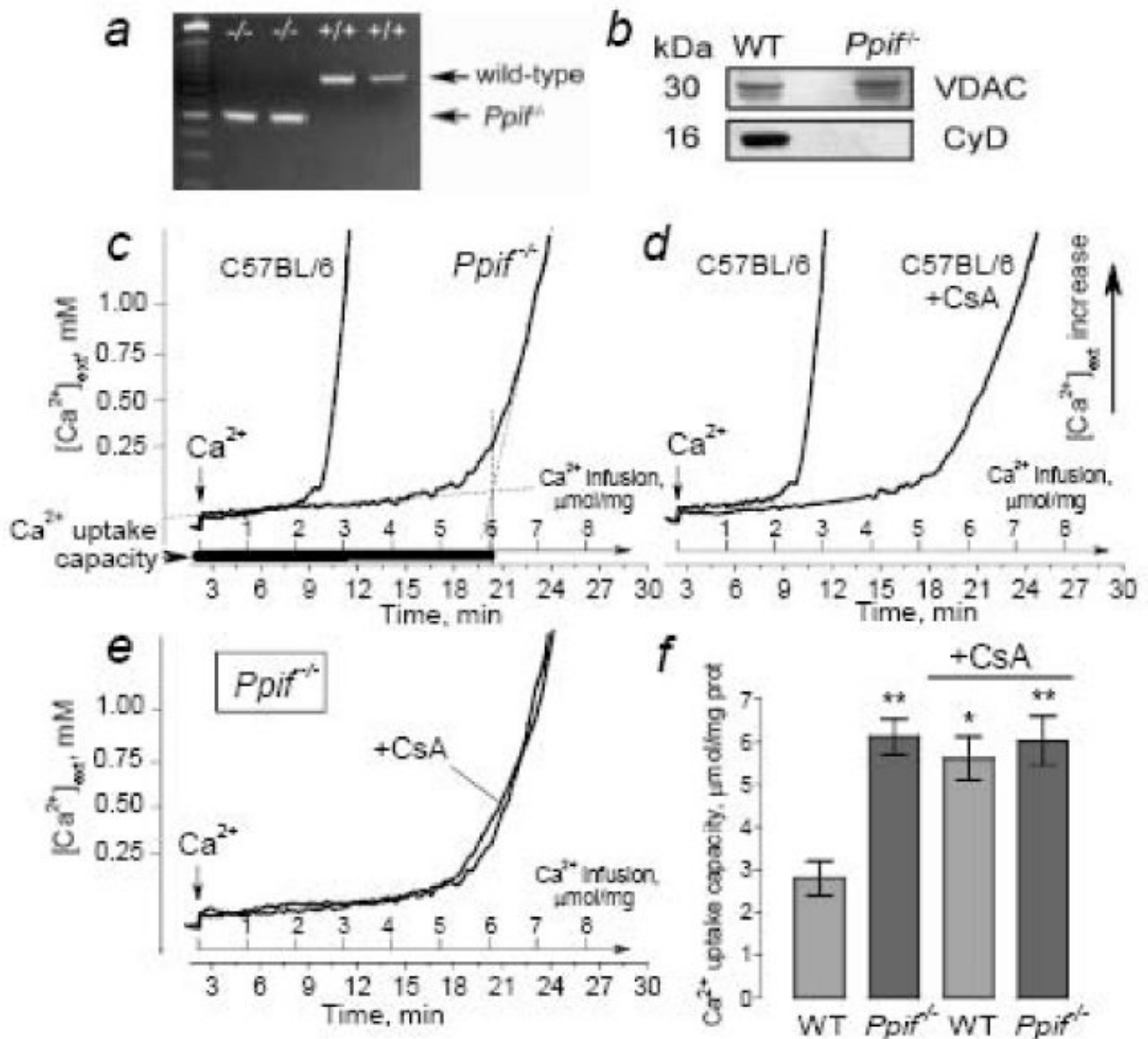
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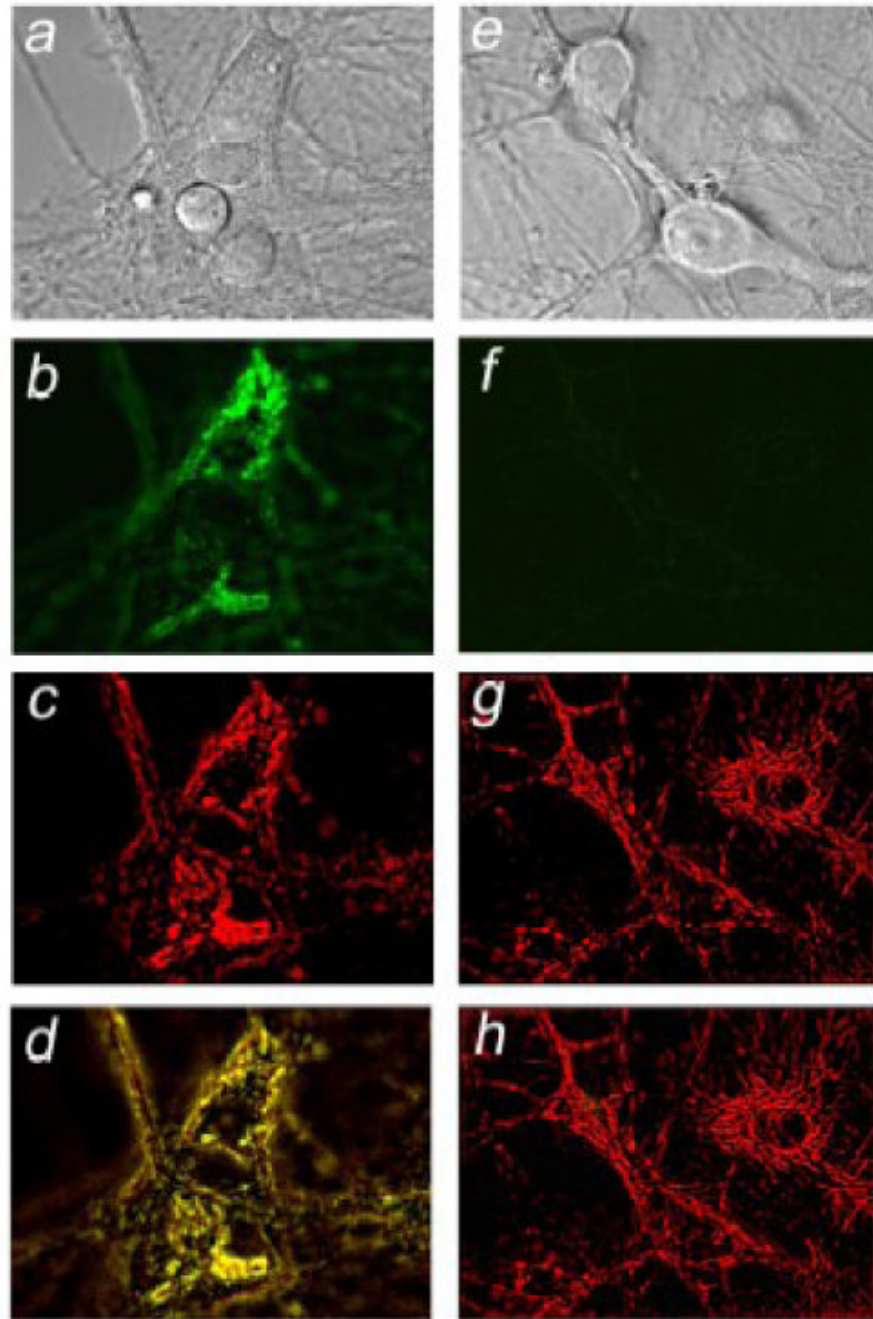




**Fig. 1.** Ca<sup>2+</sup> uptake capacity of isolated brain mitochondria from cyclophilin D-knockout mice (*Ppif*<sup>-/-</sup>) and their genetic background C57BL/6 mice (wild-type, WT). Cyclosporin A (CsA, 1 μM) augments Ca<sup>2+</sup> uptake capacity of mitochondria from C57BL/6 mice but fails to increase Ca<sup>2+</sup> uptake capacity of mitochondria from *Ppif*<sup>-/-</sup> mice

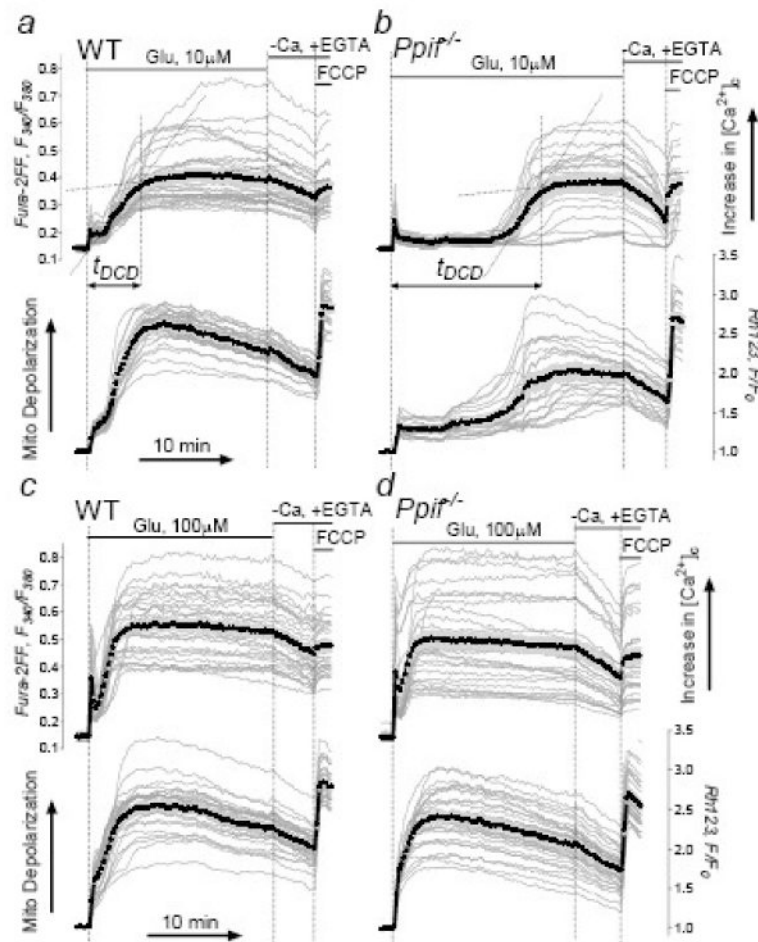
In **a**, genotyping CyD-knockout and wild-type mice: PCR products generated from genomic DNA obtained from cyclophilin D-knockout mice (*Ppif*<sup>-/-</sup>) and C57BL/6 mice (wild-type, WT) were separated by agarose gel electrophoresis and visualized with ethidium bromide. In **b**, western blot analysis of CyD expression in brain mitochondria isolated from *Ppif*<sup>-/-</sup> and wild-type (WT) mice. The protein loading for electrophoresis was 15 μg protein per lane. VDAC was used as a loading control. In **c**, comparison of Ca<sup>2+</sup> uptake capacity of brain mitochondria from C57BL/6 and *Ppif*<sup>-/-</sup> mice. In **d**, effect of CsA (1 μM) on Ca<sup>2+</sup> uptake capacity of mitochondria from C57BL/6 mice. In **e**, lack of CsA (1 μM) effect on Ca<sup>2+</sup> uptake capacity of mitochondria from *Ppif*<sup>-/-</sup> mice. In **f**, summary of data obtained with mitochondria from wild-type and *Ppif*<sup>-/-</sup> mice and incubated with and without 1 μM CsA. Data are mean ± SEM. \**p* < 0.05,

$t=4.116$ , in comparison of  $\text{Ca}^{2+}$  uptake capacity in WT mitochondria incubated with or without CsA,  $N=3$ ;  $**p<0.01$  in comparison of  $\text{Ca}^{2+}$  uptake capacity in WT mitochondria versus  $\text{Ca}^{2+}$  uptake capacity in  $Ppif^{-/-}$  mitochondria incubated with ( $t=4.742$ ) or without CsA ( $t=4.860$ ).

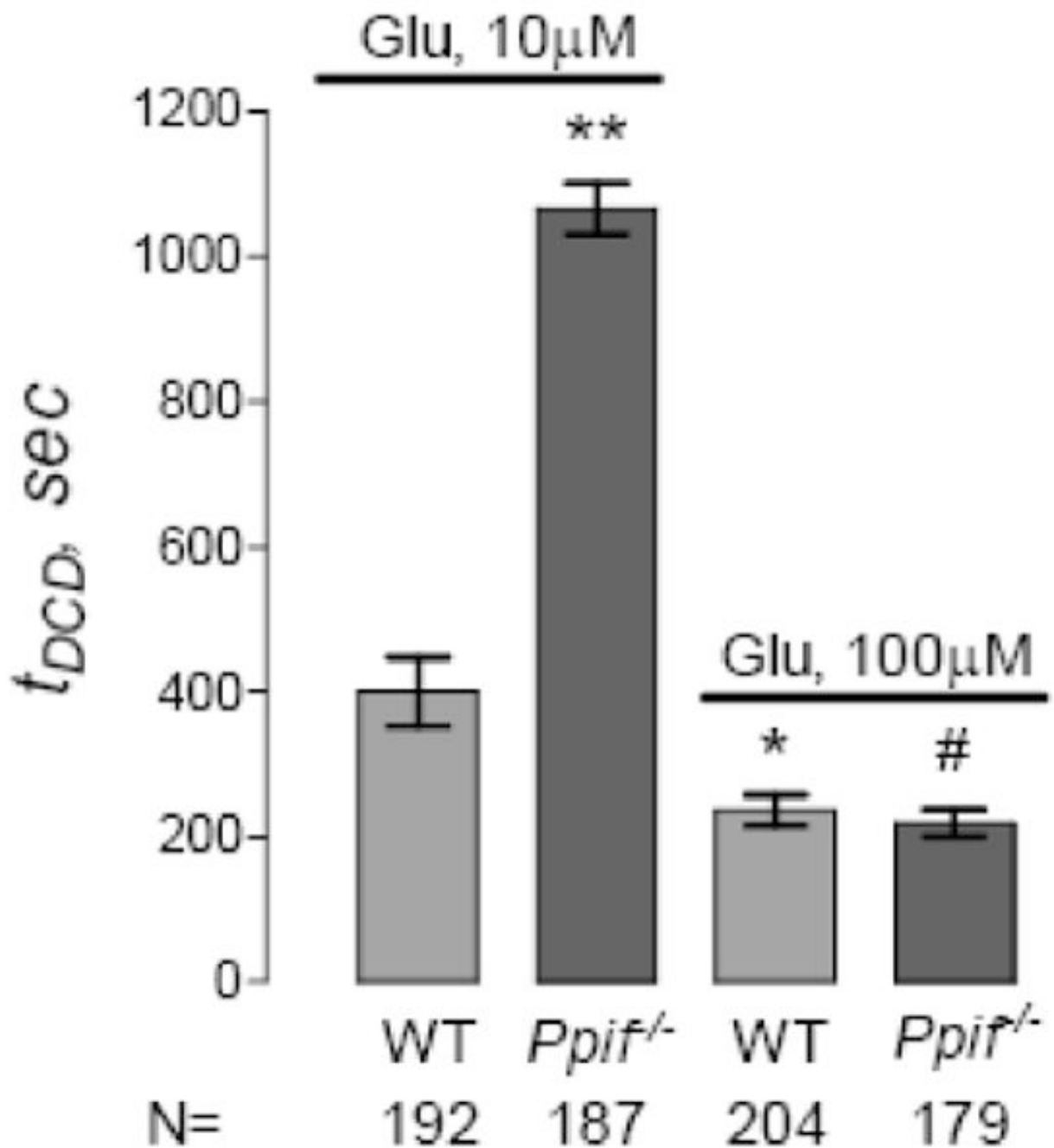


**Fig. 2. Immunocytochemical detection of cyclophilin D (CyD) in cultured cortical neurons derived from wild-type C57BL/6 mice (a-d) and cyclophilin D-knockout (*Ppif*<sup>-/-</sup>) mice (e-h)**

In *a* and *e*, phase contrast bright field images of cortical neurons from C57BL/6 and *Ppif*<sup>-/-</sup> mice, respectively. In *b* and *f*, neurons from wild-type and *Ppif*<sup>-/-</sup> mice, respectively, were exposed to mouse anti-CyD antibody and donkey anti-mouse antibody conjugated with Alexa Fluor 488 (Invitrogen). In *c* and *g*, mitochondrial staining with MitoTracker Red in neurons from C57BL/6 and *Ppif*<sup>-/-</sup> mice, respectively. In *d* and *h*, co-localization of CyD and MitoTracker staining in neurons from wild-type and *Ppif*<sup>-/-</sup> mice.



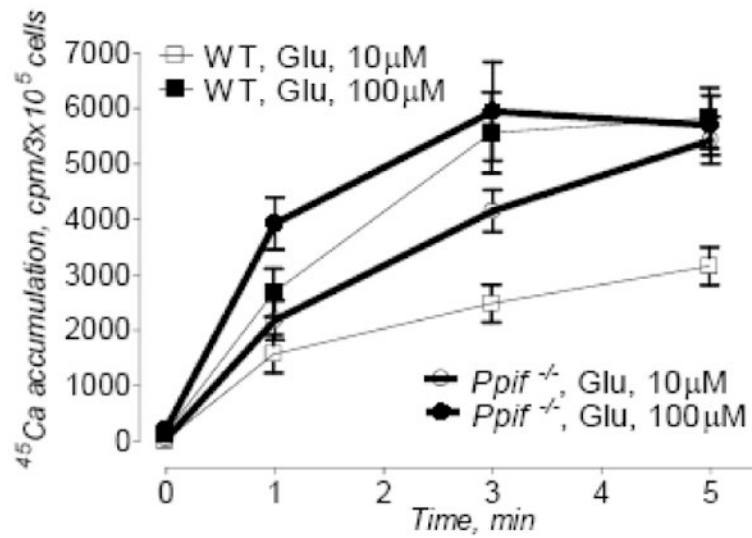
**Fig. 3. The changes in cytosolic Ca<sup>2+</sup> ([Ca<sup>2+</sup>]<sub>c</sub>) and mitochondrial membrane potential (Δψ) in response to elevated glutamate (Glu) in WT (a,c) and *Ppif*<sup>-/-</sup> (b,d) neurons**  
 In a-d, the original (thin grey traces) and the averaged fluorescence traces (thick black traces, mean ± SEM) from the representative experiments are shown. Cytosolic Ca<sup>2+</sup> was followed by monitoring Fura-2FF fluorescence and Δψ was followed by monitoring Rhodamine-123 (Rh123) fluorescence. In these experiments, neurons were exposed to 10 μM or to 100 μM glutamate plus 10 μM glycine as indicated. At the end of the experiments, glutamate and Ca<sup>2+</sup> were removed by replacing the bath solution with glutamate- and Ca<sup>2+</sup>-free solution containing 1 mM EGTA. 1 μM FCCP was added to neurons as indicated to completely depolarize mitochondria and release Ca<sup>2+</sup> accumulated in mitochondria. In a, *t*<sub>DCD</sub> is the time from the beginning of glutamate exposure to the completion of the DCD.



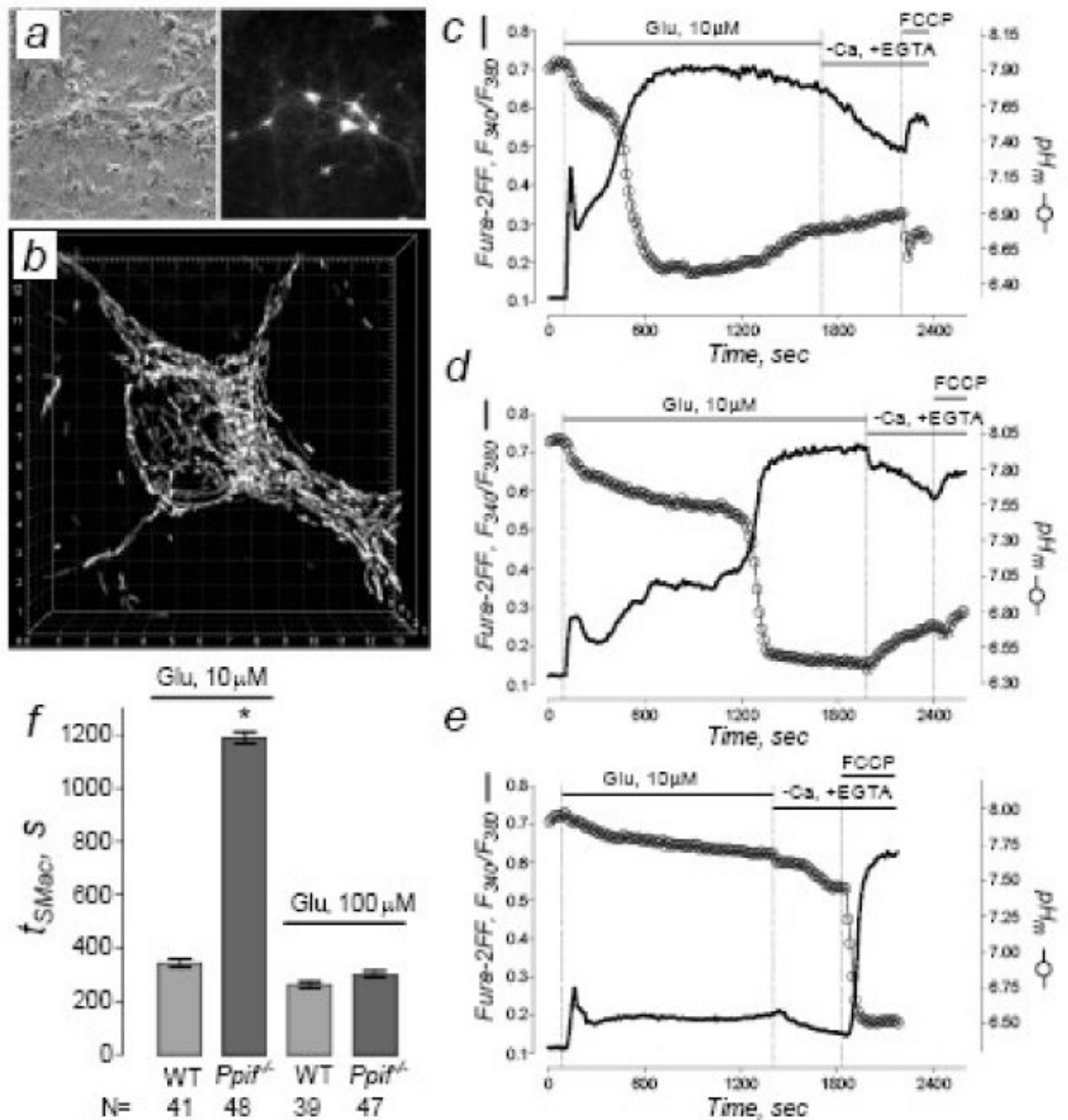
**Fig. 4. Statistical analysis of  $t_{DCD}$  obtained with WT and *Ppif*<sup>-/-</sup> neurons exposed to moderate (10 $\mu$ M) and high (100 $\mu$ M) glutamate concentrations**

Data are mean $\pm$ SEM.  $t_{DCD}$  was determined by finding the time between the beginning of glutamate exposure and the intersection point of two linear graphs approximating the uprising fragment of the averaged Fura-2FF fluorescence trace and the fragment corresponding to the elevated  $[Ca^{2+}]_c$  plateau as shown in Fig. 3a. Data are mean $\pm$ SEM. \* $p$ <0.01,  $t$ =3.525, in a comparison between WT neurons exposed to 10 and 100 $\mu$ M glutamate; \*\* $p$ <0.001,  $t$ =14.35, in a comparison between WT and *Ppif*<sup>-/-</sup> neurons exposed to 10 $\mu$ M glutamate; # $p$ <0.001,  $t$ =18.25, in a comparison between *Ppif* neurons exposed to 10 and 100 $\mu$ M glutamate. There were 8 independent experiments. N in the Figure shows the total number of cells examined in

these experiments. Statistical analysis of the experimental results consisted of one-way ANOVA followed by Bonferroni's *post hoc* test (GraphPad Prism<sup>®</sup> 4.0, GraphPad Software Inc., San Diego, CA).



**Fig. 5.**  $^{45}\text{Ca}$  accumulation in glutamate-treated WT (thick lines) and *Ppif*<sup>-/-</sup> neurons (thin lines) Neurons were incubated in the standard bath solution supplemented with 5µCi of  $^{45}\text{Ca}$ .  $^{45}\text{Ca}$  accumulation was measured at 1, 3, and 5 minutes after application of 10µM (open symbols) or 100µM glutamate (filled symbols) as indicated. Non-specific  $^{45}\text{Ca}$  binding, measured in the presence of 10µM MK801 (an inhibitor of NMDA receptors), 10µM CNQX (an inhibitor of AMPA receptors), and 5µM nifedipine (a blocker of voltage-gated  $\text{Ca}^{2+}$  channels) was subtracted from  $^{45}\text{Ca}$  accumulation in the absence of inhibitors. Data are mean±SEM, N=3.

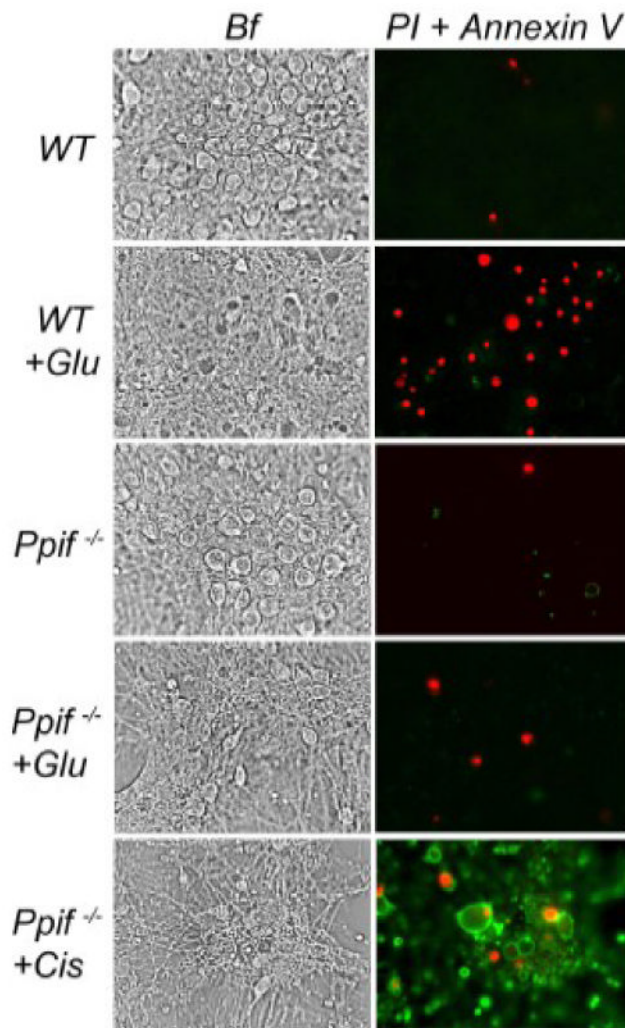


**Fig. 6. The relationship between delayed calcium deregulation and induction of the mitochondrial permeability transition in WT and *Ppif*<sup>-/-</sup> neurons**

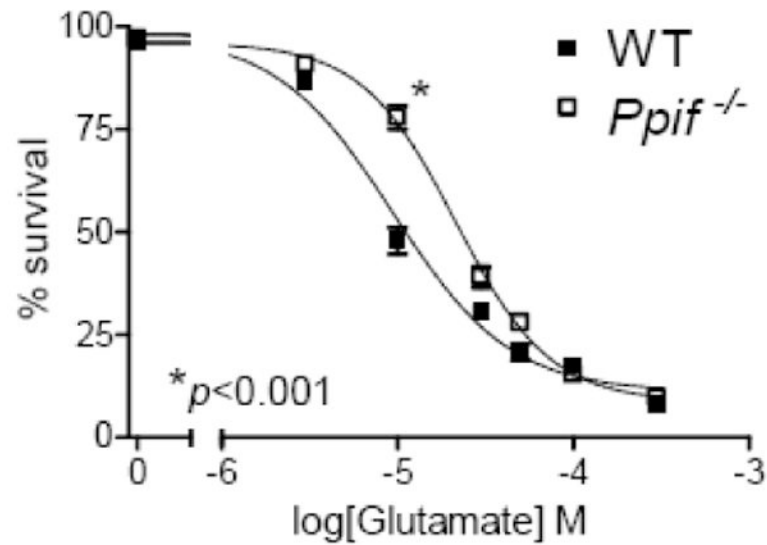
In **a**, representative bright field (right) and fluorescent (left) images of cultured hippocampal neurons expressing mito-eYFP. In **b**, a 3D reconstruction of the mitochondrial network in the neuron expressing mito-eYFP. In **c-e**, simultaneous recordings of cytosolic  $Ca^{2+}$  with Fura-2FF (*thick traces*) and matrix pH ( $pH_m$ ) with mito-eYFP (*traces with symbols*). Glutamate (Glu, 10  $\mu$ M, plus glycine, 10  $\mu$ M) was added as indicated. In **f**, statistical analysis of  $t_{SMac}$  measured with WT and *Ppif*<sup>-/-</sup> neurons exposed to moderate (10  $\mu$ M) and high (100  $\mu$ M) glutamate concentrations.  $t_{SMac}$  is the time from the beginning of glutamate exposure to the onset of the secondary matrix acidification. Data are mean  $\pm$  SEM. \* $p < 0.01$ ,  $t = 3.575$ , in a comparison



between WT neurons exposed to 10 and 100 $\mu$ M glutamate;  $**p < 0.001$ ,  $t = 37.01$ , in a comparison between WT and *Ppif*<sup>-/-</sup> neurons exposed to 10 $\mu$ M glutamate;  $#p < 0.001$ ,  $t = 38.83$ , in a comparison between *Ppif*<sup>-/-</sup> neurons exposed to 10 and 100 $\mu$ M glutamate. N in the Figure shows the total number of cells examined in these experiments. Statistical analysis of the experimental results consisted of one-way ANOVA followed by Bonferroni's *post hoc* test (GraphPad Prism<sup>®</sup> 4.0, GraphPad Software Inc., San Diego, CA).



**Fig. 7. An evaluation of necrotic cell death and the induction of apoptosis with propidium iodide (PI) and Annexin V staining after exposure of WT and *Ppif*<sup>-/-</sup> neurons to glutamate or cisplatin** Neurons were exposed for 10 minutes to 10 $\mu$ M glutamate plus 10 $\mu$ M glycine (+Glu) as indicated. Then, glutamate was removed and necrotic cell death or induction of apoptosis was evaluated 24 hours later following the appearance of staining with PI or Annexin V. *Ppif*<sup>-/-</sup> neurons treated with cisplatin (+Cis, 30 $\mu$ M, 24 hours) were used as a positive control for Annexin V staining. *Bf*, phase contrast bright field. These experiments with similar results were performed in triplicate with neurons from three different platings.



**Fig. 8. Dose-dependence of glutamate-induced neuronal death**

Trypan Blue exclusion method was used to evaluate neuronal death (Dubinsky and Rothman, 1991). Neurons derived from wild-type (WT) C57BL/6 and *Ppif*<sup>-/-</sup> mice were exposed to various glutamate concentrations (3-300 $\mu$ M) for 10 minutes, then glutamate was removed and cell death was evaluated after 24 hours by counting Trypan Blue stained neurons in a blind manner. Data are mean $\pm$ SEM. \* $p < 0.001$ ,  $t = 12.12$ , in a comparison between WT and *Ppif*<sup>-/-</sup> neurons exposed to 10 $\mu$ M glutamate. N=9.