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Promiscuous actions of small molecule inhibitors of the protein kinase D-class IIa HDAC axis in striated muscle



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1. Introduction

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ABSTRACT

PKD-mediated phosphorylation of class IIa HDACs frees the MEF2 transcription factor to activate genes that govern muscle differentiation and growth. Studies of the regulation and function of this signaling axis have involved MC1568 and Gö-6976, which are small molecule inhibitors of class IIa HDAC and PKD catalytic activity, respectively. We describe unanticipated effects of these compounds. MC1568 failed to inhibit class IIa HDAC catalytic activity in vitro, and exerted divergent effects on skeletal muscle differentiation compared to a bona fide inhibitor of these HDACs. In cardiomyocytes, Gö-6976 triggered calcium signaling and activated stress-inducible kinases. Based on these findings, caution is warranted when employing MC1568 and Gö-6976 as pharmacological tool compounds to assess functions of class IIa HDACs and PKD.

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The 18 mammalian histone deacetylases (HDACs) are grouped into four classes (I, II, III and IV) [1]. Class II HDACs are further divided into two subclasses, IIa (HDACs 4, 5, 7 and 9) and IIb (HDACs 6 and 10). Class IIa HDACs serve critical roles in the control of muscle gene expression [2]. These HDACs have long (~500 amino acid) amino-terminal extensions that harbor binding sites for pro-myogenic transcription factors such as myocyte enhancer

for pro-myogenic transcription factors such as myocyte enhancer factor 2 (MEF2) [3–7]. In skeletal muscle cells, binding of class IIa HDACs to MEF2 results in suppression of MEF2 target genes that govern differentiation of myoblasts into multi-nucleated myotubes [8,9]. In cardiac muscle, interaction of class II HDACs with MEF2 blocks hypertrophic growth of the heart [10].

The amino-terminal extensions of class IIa HDACs also harbor two serine residues that confer signal-responsiveness to the enzymes. These sites are phosphorylated by members of the calcium/calmodulin-dependent kinase (CaM kinase) superfamily,

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which includes CaM kinase II and protein kinase D (PKD) [11– 14]. Phosphorylation of class IIa HDACs creates docking sites for the 14-3-3 chaperone protein, which promotes nuclear export of the transcriptional repressors. Thus, in response to CaMK/PKD signaling, class IIa HDACs dissociate from MEF2, freeing the transcription factor to activate downstream target genes. The identification of PKD as a class IIa HDAC kinase was aided by the use of a pharmacological tool compound, Gö-6976 [15], which functions as an ATP-competitive inhibitor of the three PKD isoforms, PKD-1, -2 and -3 [16].

Despite having conserved catalytic domains, class IIa HDACs are unable to deacetylate histones [17]. In fact, bona fide substrates of class IIa HDACs have yet to be identified, and class IIa HDAC catalytic activity can only be monitored using an artificial substrate [18]. Class IIa HDAC catalytic activity is largely resistant to commonly used HDAC inhibitors, such as suberanilohydroxamic acid (SAHA) [19]. Several years ago, MC1568 (originally referred to as compound 2f) was described as the first class IIa HDAC-selective inhibitor [20]. MC1568 has been used in a multitude of studies as a chemical biological probe to assess the function of class IIa HDAC catalytic activity in various physiological and pathophysiological processes. For example, MC1568 was recently found to block pulmonary artery endothelial cell proliferation and rescue experimental pulmonary hypertension [21], as well as block hepatic stellate cell activation and liver fibrosis in mice [22].

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Abbreviations: HDAC, histone deacetylase; PKD, protein kinase D

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Paradoxically, MC1568 was shown to block skeletal muscle differentiation by stabilizing MEF2:HDAC4 complexes and preventing MEF2 acetylation [23].

In the current studies, we confirm that MC1568 blocks differentiation of C2C12 myoblasts into multinucleate myotubes. However, the anti-myogenic activity of this compound appears to be due to off-target action(s), since enzymatic assays revealed an inability of MC1568 to inhibit class IIa HDAC catalytic activity. Furthermore, we validate that Gö-6976 is a PKD inhibitor. Surprisingly, however, exposure of primary cardiomyocytes to Gö-6976 leads to enhanced calcium signaling and activation of stress-inducible kinases such as p38 mitogen-activated protein kinase and c-Jun N-terminal kinase (JNK). These findings highlight the need to employ multiple, structurally distinct compounds when assessing the functions of class IIa HDACs and PKD using pharmacological tools.

2. Materials and methods

2.1. Inhibitors, antibodies and adenoviruses

Gö-6976 was obtained from EMD Millipore and A.G. Scientific. MC1568 was purchased from both Sigma-Aldrich and Selleck Chemicals. SAHA, MGCD0103 and Tubastatin A were purchased from Selleck Chemicals, whereas diphenylacetohydroxamic acid (DPAH) [24] was obtained from Sigma-Aldrich (D6071). DPAH was also synthesized in-house, and its purity was confirmed to be greater than 95%. U-73122 was procured from Tocris Bioscience. For all studies, DMSO was used as the compound vehicle (final DMSO concentration = 0.1%). For immunoblotting, cells were homogenized in PBS (pH 7.4) containing 0.5% Triton X-100, 300 mM NaCl and protease/phosphatase inhibitor cocktail (Thermo Fisher) using a Bullet Blender homogenizer (Next Advance). The following. Homogenates were sonicated and clarified by centrifugation at $12000 \times g$ for 10 minutes. Antibodies were used for immunoblotting: anti-calnexin (Santa Cruz Biotechnology; sc-11397), anti-PKD1 (Cell Signaling Technology; #2052), anti-phospho-PKD1 Ser-738/742 (Cell Signaling Technology; #2054), anti-phospho-PKD1 Ser-910 (Cell Signaling Technology; #2051), anti-phospho-ERK1/2 (Cell Signaling Technology; 4370), anti-phospho-p38 (Cell Signaling Technology; 4631) and anti-phospho-JNK (Cell Signaling Technology; 4668). Adenoviruses for wild-type PKD1 and catalytically inactive PKD1 harboring a K612W substitution (K/W) [25] were generated using pShuttle-CMV, as previously described [26].

2.2. Primary cardiac myocyte preparation

Neonatal rat ventricular myocytes (NRVMs) were prepared from hearts of 1- to 3-day-old Sprague-Dawley rats, as previously described [27]. Cells were cultured overnight on 10-cm plates coated with gelatin (0.2%; Sigma-Aldrich) in Dulbecco's Modified Eagle's Medium (DMEM) containing calf serum (10%), L-glutamine (2 mM), and penicillin-streptomycin. After overnight culture, cells were washed with serum-free medium and maintained in DMEM supplemented with L-glutamine, penicillin-streptomycin, and Neutridoma-SP (0.1%; Roche Applied Science), which contains albumin, insulin, transferrin, and other defined organic and inorganic compounds. Adult rat ventricular myocytes (ARVMs) were obtained from female Sprague-Dawley rats, as described previously [28]. ARVMs were plated at a density of 100-150 cells/ mm² on laminin-coated 60-mm plates and maintained in serumfree DMEM supplemented with albumin (2 mg/mL), 2,3-butanedione monoxime (1 mg/mL), L-carnitine (2 mM), creatine (5 mM), penicillin-streptomycin (100 µg/mL), triiodothyronine (1 pM), and taurine (5 mM).

2.3. C2C12 culture and transfection

C2C12 myoblasts were obtained from ATCC and were maintained in DMEM containing 10% FBS, L-glutamine (2 mM), and penicillin–streptomycin. C2C12 cells were induced to differentiate by replacing FBS in the culture medium with 2% horse serum. C2C12 cells were transfected with a $3 \times$ MEF2-luciferase plasmid, which contains three tandem copies of the mouse muscle creatine kinase enhancer MEF2 site upstream of the thymidine kinase promoter [29], using FuGENE 6 (Promega). To control for non-specific effects of compounds on luciferase expression, parallel plates of C2C12 cells were transfected with a TATA-luciferase plasmid lacking MEF2 binding sites. Luciferase activity is expressed as a ratio of MEF2-luciferase:TATA-luciferase.

2.4. In vitro kinase assays

NRVMs on 10-cm plates (2×10^6 cells/plate) were serum starved for 4 h and subsequently exposed to phenylephrine $(10 \,\mu\text{M})$ for 1 h. Following treatment, total protein lysates were prepared, as described above for immunoblotting. Total protein (200 µg) was incubated in buffer (500 µl) containing antibodies specific for PKD1 (Santa Cruz Biotechnology; sc-639), PKD2 (Bethyl Laboratories; BL844) or PKD3 (Bethyl Laboratories; BL1282), overnight at 4 °C with rocking. Antibody-protein complexes were captured with Protein G-Sepharose beads (GE Healthcare Life Sciences), with incubation for 1 h at 4 °C. Immunoprecipitates were equilibrated with kinase buffer (20 mM HEPES, pH 7.5, 20 mM MgCl₂), and kinase reactions were initiated upon addition of ATP (0.1 mM), 10 μ Ci [γ -³²P]-ATP, and 2 µg Syntide-2 substrate (Anaspec). Kinase reactions were carried out at 30 °C for 30 min, and terminated with 2× SDS-PAGE loading dye. Phospho-Syntide-2 peptide was resolved by SDS-PAGE and visualized by autoradiography.

2.5. Calcium measurements

NRVMs were loaded with 5 µM Fluo 3-AM and 0.05% pluronic acid (both from Invitrogen) dissolved in Tvrode's solution (in mM: 140 NaCI, 5 KCl, 1 MgCl₂, 1 CaCl₂, 10 HEPES, 5 Glucose, pH adjusted to 7.4 with NaOH) for 20 min. NRVMs were then washed three times in Tyrode's solution with gentle agitation. After a 10 min de-esterification period, Fluo-3 AM-loaded cells, bathed in Tyrode's solution, were placed on the stage of an LSM 510 META scanning laser confocal microscope, and viewed with a Zeiss Plan-Neofluar $40 \times$ objective (Zeiss). Fluo-3 AM was excited with the 488-nm line of an argon laser (30-mW maximum output, operated at 50% or 6.3A, attenuated to 5%). The emitted fluorescence was directed through a dual 488/543 dichroic mirror to a photomultiplier equipped with a 505 nm long-pass filter. Caffeine (Sigma-Aldrich) or Gö-6976 (Sigma-Aldrich) were diluted to working concentrations (5 mM and 1 µM, respectively) in Tyrode's solution, and were delivered via a manually-operated, gravity-driven global perfusion system. Spontaneous Ca²⁺ fluctuations were analyzed qualitatively using NIH Image J software. Fluorescence amplitude data are expressed as $\Delta F/F$, where F represents baseline fluorescence prior to application of caffeine or Gö-6976, and ΔF represents the change in peak fluorescence during the application of either compound. All Ca²⁺ imaging experiments were performed at room temperature (~25 °C) in the University of Colorado Denver Advanced Light Microscopy Core.

2.6. Immunostaining

Following experimental treatments, NRVMs were washed twice in PBS (pH 7.4), fixed with formalin (10%) in PBS, permeabilized and blocked with PBS containing NP-40 (0.1%) and bovine serum albumin (BSA; 3%), and then incubated in the same solution containing primary antibody specific for PKD1 phosphorylated on serine-916 (1:1000 dilution; Cell Signaling Technology; #2051) overnight at 4 °C. Cells were washed four times in PBS and incubated in PBS-NP-40-BSA containing a Cy3-conjugated secondary antibody (1:200 dilution; Jackson Laboratories) for 30 min at room temperature. Cells were washed four times in PBS and then covered with mounting solution (SlowFade; Molecular Probes) and glass coverslips. For myosin staining of skeletal muscle cells, immunohistochemical analysis was performed using an anti-sarcomeric myosin (MF20; Iowa Hybridoma Bank) primary antibody, a secondary antibody (biotinylated anti-mouse IgG; Vector Labs), and Elite ABC and VECTOR NovaRED Substrate Kits (Vector Labs). The percentage of the plate containing myosin-positive myotubes was quantified in a blinded manner using an Axiovert 200 inverted microscope with a digital camera equipped with AxioVision imaging software (Zeiss). Values were averaged from five independent plates of cells per condition.

2.7. HDAC activity assays

HDAC activity assays were completed as previously reported [30]. Each HDAC substrate was based on ε -N-acylated lysine, derivatized on the carboxyl group with amino-methylcoumarin (AMC) [18]. Subsequent to deacylation by HDACs, trypsin was used to release AMC, resulting in a significant increase in fluorescence. Rat heart homogenates were employed as a source of HDAC activity. Homogenates were prepared in PBS (pH 7.4) containing 0.5% Triton X-100, 300 mM NaCl and protease/phosphatase inhibitor cocktail (Thermo Fisher), using a Bullet Blender homogenizer (Next Advance). Tissue extracts were diluted into PBS in 100 µl total volume in a 96-well plate (60 µg ventricular protein/well). Tissue extracts were incubated with increasing concentrations of HDAC inhibitors for 1 h. Substrates were added (5 µl of 1 mM DMSO stock solutions), and the plates were placed at 37 °C for 2–3 h. Finally, developer/stop solution was added (50 µl per well of PBS with 1.5% Triton X-100, 3 µM TSA, and 0.75 mg/mL trypsin). followed by an additional 20 min incubation at 37 °C. AMC fluorescence was measured using a BioTek Synergy 2 plate reader, with excitation and emission filters of 360 nm and 460 nm, respectively (each with 40 nm bandwidth), along with a 400 nm dichroic top mirror. As a control, HDAC inhibitors were incubated with AMC without the appended substrate. The importance of this step is illustrated by the fact that MC1568 quenches AMC-generated fluorescence (sFig. 1). Background signals from buffer blanks were subtracted, and the data were normalized to appropriate controls. To monitor recombinant HDAC4 activity, 15 ng of recombinant human HDAC4 (BPS Bioscience; #50004) or PBS blank was added to a 96-well plate. Wells were incubated for 1 h with increasing concentrations of DPAH or MC1568 (Sigma-Aldrich), prior to addition of the cell-permeable class IIa fluorescent HDAC substrate. After two hours, stop buffer was added to each well and incubated for 20 min at 37 °C, and deacetylase activity was quantified as described above.

3. Results

3.1. Differential effects of class IIa HDAC inhibitors on skeletal muscle differentiation

Association of MEF2 with class IIa HDACs leads to repression of downstream target genes that promote skeletal muscle differentiation (Fig. 1A). When class IIa HDACs are phosphorylated by kinases such as PKD, these HDACs are exported to the cytoplasm, and MEF2 becomes transcriptionally active. Gö-6976 inhibits the catalytic activity of PKD, while MC1568 and DPAH are small molecule inhibitors of class IIa HDAC catalytic activity (Fig. 1A and B).

To assess the impact of isoform-selective HDAC inhibition on skeletal muscle differentiation, C2C12 myoblasts were differentiated in the absence or presence SAHA (pan-HDAC inhibitor), MGCD0103 (class I HDAC inhibitor), Tubastatin A (HDAC6 inhibitor), MC1568 or DPAH. Pre-treatment with SAHA and MGCD0103 led to complete inhibition of C2C12 cell differentiation, as indicated by the absence of myosin-positive myotubes in these cultures (Fig. 1C). Consistent with prior findings, HDAC6 inhibition with Tubastatin A moderately impaired myogenesis [31], while the class IIa HDAC inhibitor MC1568 completely abolished formation of multinucleated myotubes [23]. Surprisingly, however, the structurally distinct class IIa HDAC inhibitor, DPAH, failed to block differentiation of C2C12 cells. Higher magnification images of cells from a second experiment with MC1568 and DPAH revealed that DPAH actually enhanced C2C12 mvotube formation (Fig. 1D and E). The differential actions of the two compounds on muscle differentiation correlated with their effects on MEF2 transcriptional activity (Fig. 1F), with MC1568 and DPAH repressing and stimulating a MEF2-dependent reporter gene, respectively.

3.2. Commercially available MC1568 does not inhibit class IIa HDAC catalytic activity

To begin to address the discrepant results with MC1568 and DPAH, the compounds were tested for their ability to block HDAC catalytic activity in vitro. Dose-response studies were performed with substrates that enable specific assessment of class I, IIa or IIb HDAC activity. Rat left ventricular homogenates were used as a rich source of endogenous HDACs [30]. Consistent with prior findings [20], MC1568 had no observable effect on class I HDAC activity (Fig. 2A). Surprisingly, however, MC1568 also failed to inhibit class IIa HDAC activity and, instead, modestly suppressed class IIb HDAC catalytic activity (Fig. 2B and C); HDAC6 is the predominant class IIb HDAC in these lysates [30]. In contrast, DPAH dose-dependently inhibited class IIa HDACs ($IC_{50} = 1.2 \mu M$), while having no effect on class I or IIb HDACs (Fig. 2D-F). Similar results were obtained using independent sources of MC1568 and DPAH (Fig. 2G-I and data not shown). To confirm these findings, recombinant HDAC4 was used as a representative class IIa HDAC. Consistent with the results obtained using endogenous HDACs, MC1568 failed to block the activity of recombinant HDAC4 (Fig. 2]), while DPAH efficiently inhibited this class IIa HDAC (Fig. 2K). Quantification of class IIa HDAC activity was enabled by the use of a substrate having a trifluoroacetyl group on the ε amine, which is deacetylated by class IIa HDACs but not class I or IIb HDACs [32].

The mechanism underlying the inability of MC1568 to inhibit class IIa HDACs could be related to compound structure. Indeed, nuclear magnetic resonance (NMR) data revealed the commercial compound to be an isomer of the previously published structure (Fig. 2L).

3.3. Gö-6976 activates stress-inducible kinases in cardiac myocytes

Among the downstream effectors of $G_{\alpha}q$ -coupled receptors in cardiac myocytes are members of the PKD family [26,33]. The three PKD isoforms exhibit a high degree of sequence similarity, consisting of ~900 amino acids, with two amino-terminal cysteine-rich domains (CRDs), an internal pleckstrin homology domain (PH), and carboxy-terminal catalytic domains [34,35]. PKD can be activated by protein kinase C (PKC)-mediated phosphorylation of activation loop sites within the catalytic domain, with subsequent auto-phosphorylation of PKD on serine-916 (Fig. 3A).



Fig. 1. Differential effects of isoform-selective HDAC inhibitors on skeletal muscle differentation. (A) A schematic representation of the PKD-class IIa HDAC axis. Gö-6976 blocks the catalytic activity of PKD, while MC1568 and DPAH inhibit the enzymatic activity of class IIa HDACs. (B) The molecular structures of the compounds used in this study are shown. (C) C2C12 myoblasts were induced to differentiate in the absence (vehicle) or presence of the indicated HDAC inhibitors; SAHA (1 μ M), MGCD0103 (500 nM), Tubastatin A (1 μ M), MC1568 (10 μ M) and DPAH (10 μ M). After 4 days of differentiation, cells were fixed and stained with an anti-myosin antibody to reveal myosin heavy chain (MyHC)-positive skeletal myotubes. (D) C2C12 cells were induced to differentiate in the absence or presence of the class IIa HDAC inhibitors MC1568 and DPAH, stained as described for (C), and differentiation was quantified (E); Scale bar = 10 μ M. (F) C2C12 myoblasts were transfected with a MEF2-dependent luciferase reporter gene, and the cells were subsequently induced to differentiate for three days in the absence or presence of MC1568 or DPAH. For (E) and (F), five independent plates of cells were analyzed per condition. Numbers represent averages + S.E.M.; *P < 0.05 vs. vehicle-treated cells.

PKD1, -2 and -3 were immunoprecipitated from NRVMs that had been stimulated with the α_1 -adrenergic receptor agonist phenylephrine (PE), which potently activates PKD in cardiac myocytes. Immunoprecipitates were incorporated into in vitro kinase reactions containing syntide-2 substrate and [γ^{-32} P]-ATP, in the absence or presence of Gö-6976. As shown in Fig. 3B, Gö-6976 effectively inhibited the ability of PKD1, -2 and -3 to phosphorylate the synthetic substrate, and also blocked auto-phosphorylation of PKD1 and PKD3; PKD2 auto-phosphorylation was not detected in the assay.

Next, the impact of Gö-6976 on auto-phosphorylation of PKD1 on serine-916 in intact, unstimulated cardiomyocytes was assessed. Paradoxically, treatment of NRVMs with this compound led to enhanced serine-916 phosphorylation, while a related compound, Gö-6983, which targets PKC but not PKD, had no effect on PKD1 phosphorylation (Fig. 3C). Stimulation of PKD1 serine-916 phosphorylation by Gö-6976 in NRVMs was confirmed by immunoblotting, and was shown to occur with compounds obtained from two independent vendors (Fig. 3D; left-hand panels). Gö-6976 similarly promoted phosphorylation of the activation loop sites on PKD1 (serines-744 and -748) in NRVMs. Furthermore, Gö-6976 was also able to increase PKD1 phosphorylation in adult rat ventricular myocytes (ARVMs) (Fig. 3D; right-hand panels), but not in HEK293 fibroblasts (Fig. 3E), suggesting cell type-specificity of the effects.



Fig. 2. MC1568 does not block class IIa HDAC catalytic activity. Dose-response studies were performed to assess the ability of MC1568 and DPAH to inhibit HDAC catalytic activity in vitro. (A–C) MC1568 (from Sigma–Aldrich) failed to inhibit class I and class IIa HDAC activity, but did block class IIb activity at higher concentrations. (D–F) DPAH dose-dependently, and selectively, inhibited class IIa HDAC activity. MC1568 from Selleck Chemicals also failed to inhibit endogenous class IIa HDAC activity (G–I). MC1568 was unable to inhibit the catalytic activity of a recombinant form of a prototypical class IIa HDAC, HDAC4, while DPAH efficiently inhibited recombinant HDAC4 (J and K). (L) Analysis of NMR data from Sigma and Selleck revealed the commercial MC1568 to be a 2,5-disubstituted pyrrole, as opposed to the 2,4-disubstituted isomer shown in Fig. 1B.

Stimulation of PKD1 serine-916 phosphorylation by Gö-6976 is paradoxical, since serine-916 is thought to be an auto-phosphorylation site, and Gö-6976 is an ATP-competitive inhibitor of PKD [16]. To determine if Gö-6976 triggers PKD auto-phosphorylation, experiments were performed with ectopic PKD1 expressed in NRVMs via adenoviruses. Consistent with the results with endogenous PKD1, ectopically expressed wild-type PKD1 was efficiently phosphorylated on serine-916 upon treatment with PE or Gö-6976 (Fig. 3F). When normalized to total PKD1 expression, serine-916 phosphorylation of catalytically inactive PKD1 (K/W) was attenuated in PE-treated cells (Fig. 3F, lane 5), consistent with α_1 -adrenergic receptor signaling causing PKD1 auto-phosphorylation [26,33]. Remarkably, however, serine-916 of PKD1 (K/W) was still phosphorylated following treatment with Gö-6976 (Fig. 3F, lane 6), suggesting that the compound stimulates PKD1 serine-916 phosphorylation through a PKD-independent mechanism governed by a distinct kinase(s). This explains how Gö-6976 can inhibit PKD catalytic activity (Fig. 3B), and still trigger PKD Ser-916 phosphorylation. Additional immunoblotting studies revealed that Gö-6976 also stimulated phosphorylation of the stress-inducible MAP kinases p38 and JNK in cardiomyocytes, but not extracellular signal-regulated kinase (ERK1/2) (Fig. 3G).



Fig. 3. Gö-6976 activates stress-dependent kinases in cardiomyocytes. (A) Schematic representation of PKD1, with cysteine-rich domains (CRD) and a pleckstrin homology (PH) domain indicated. PKC phosphorylates activation loop sites in the catalytic domain of PKD1 (serines-744 and -748), resulting in stimulation of the kinase. Serine-916 can undergo auto-phosphorylation. (B) NRVMs were stimulated with phenylephrine (PE; 10 µM) for 1 h. Whole cell lysates were immunoprecipitated (IP) with antibodies to PKD1, -2 or -3 and incorporated into in vitro kinase reactions in the absence or presence of Gö-6976 (10 µM). Gö-6976 (nibited the ability of each PKD isoform to phosphorylate the synthetic syntide substrate, and blocked auto-phosphorylation of PKD1 and PKD3. (C) NRVMs were treated with Gö-6976 (10 µM) or Gö-6983 (10 µM) for 1 h, and PKD serine-916 phosphorylation was assessed by indirect immunofluorescence. Scale bar = 10 µM. (D) NRVMs (left panel) or adult rat ventricular myocytes (right panel) were treated for 1 h with vehicle control or Gö-6976 (10 µM) from two independent commercial sources, as described in the Section 2. Protein homogenates were immunoblotted with antibodies to the indicated forms of PKD1. (E) HEK293A fibroblasts were treated for 1 h with Gö-6976 (10 µM) or phorbol-12-myristate-13-acetate (PMA; 50 nM), which served as a positive control for PKC and PKD activation. Immunblotting was performed as in (D). (F) NRVMs were infected with adenoviruses encoding wild-type PKD1 or PKD1K/W, which is catalytically inactive. Cells were treated for 1 h with PE (10 µM) or Gö-6976 (10 µM), lysed, and immunoblotted with antibodies to the indicated for 1 h with Gö-6976 (10 µM) from two different vendors, and protein homogenates were immunoblotted with antibodies to the indicated for 1 h with Gö-6976 (10 µM) or Gö-6976 (10 µM), lysed, and immunoblotted with the indicated proteins.

3.4. Gö-6976 alters calcium signaling in cardiac myocytes

Given the ability of Gö-6976 to activate multiple kinases (PKC, p38 and JNK) in cultured cardiomyocytes, experiments were next performed to address the possibility that the compound functions proximally, via a general signaling mediator(s), to stimulate diverse downstream events. Consistent with this notion, the

phospholipase C (PLC) inhibitor U-73122 blocked Gö-6976mediated PKD phosphorylation as efficiently as it blocked PEinduced phosphorylation of the kinase (Fig. 4A). Subsequently, effects of Gö-6976 and caffeine on calcium signaling in cardiomyocytes were assessed by imaging cells that were loaded with a calcium-sensitive dye, Fluo-3 AM. Spontaneous calcium transients were detected in NRVMs prior to application of either caffeine or Gö-6976 (Fig. 4B). Caffeine treatment of NRVMs led to induction of sarcoplasmic reticulum (SR) calcium release (Fig. 4B, top panel). Strikingly, Gö-6976 also triggered myoplasmic calcium fluctuations within seconds after addition to NRVMs (Fig. 4B, bottom panel). Effects of caffeine and Gö-6976 on NRVM calcium were distinct, with Gö-6976-induced transients having lesser amplitude than spontaneous transients. Furthermore, Gö-6976 treatment caused a rapid increase in myoplasmic calcium stores.

4. Discussion

The current study demonstrates that MC1568 and Gö-6976 exert paradoxical effects when delivered to striated muscle cell



Fig. 4. Gö-6976 alters calcium signaling in cardiac myocytes. (A) NRVMs were pretreated with the phospholipase C (PLC) inhibitor U-73122 (1 μ M) for 20 min prior to addition of phenylephrine (PE; 10 μ M) or Gö-6976 (1 or 10 μ M) for 1 h. Immunoblotting was performed to assess levels of phospho- and total PKD1. (B) NRVMs were loaded with the calcium-sensitive dye, Fluo-3 AM. Calcium fluctuations prior to and after treatment with caffeine (5 mM) or Gö-6976 (1 μ M) were measured using a confocal microscope. Fluorescence amplitude data are expressed as $\Delta F/F$, where *F* represents baseline fluorescence prior to application of caffeine or Gö-6976, and ΔF represents the change in peak fluorescence during the application of either compound.

cultures. Consistent with prior results, MC1568 inhibited differentiation of C2C12 myoblasts into multinucleate myotubes. However, this compound failed to block the enzymatic activity of class IIa HDACs in in vitro assays. Identical findings were made with MC1568 acquired from two different vendors. In contrast to the data obtained with MC1568, a structurally distinct compound, DPAH, dose-dependently inhibited class IIa HDAC catalytic activity and enhanced, rather than inhibited, C2C12 differentiation. The contrapositive effects of the compounds on muscle differentiation correlated with effects on MEF2 transcriptional activity, with MC1568 and DPAH inhibiting and activating a MEF2-driven reporter gene, respectively. These results suggest that MC1568 inhibits skeletal myogenesis through a mechanism that is independent of suppression of class IIa HDAC catalytic activity, perhaps through an off-target action. Furthermore, the findings with DPAH illustrate that inhibition of class IIa HDAC catalytic activity is not sufficient to block myogenesis, as previously suggested, and, conversely, may enhance muscle differentiation.

At least three explanations could account for our inability to replicate the previous finding that MC1568 inhibits class IIa HDAC catalytic activity. First, analysis of NMR spectra revealed that commercially available MC1568 is an isomer of the published structure. Of note, a recent medicinal chemistry report of the resynthesis of MC1568 defined the major synthetic product to be the same isomer sold by Sigma and Selleck [36]. As such, we cannot rule out the possibility that the differential effects on class IIa HDAC activity are due to distinct properties of the isomers. A second possibility is related to the assays that were used to define class IIa HDAC catalytic activity. We monitored catalytic activity of the HDACs with a synthetic substrate that is specifically deacetylated by class IIa HDACs; this substrate is not deacetylated by other HDAC isoforms. In contrast, Mai et al. used a histone H4-³H-acetyl-peptide substrate [20]. However, it should be noted that follow-up studies by Mai and others did employ the same AMC-based fluorescent class IIa substrate used in our work [23], and we have found that the AMC fluorophore is non-specifically quenched by MC1568 (sFig. 1). Thus, a third explanation for our disparate findings is that MC1568 creates a false positive reading by quenching AMC-generated fluorescence of non-histone synthetic substrates. We circumvented this issue by normalizing the signal obtained upon addition of MC1568 to AMC-class IIa HDAC substrate to the signal obtained with AMC alone (Fig. 2B, H and I; see Materials and methods).

We also show that the staurosporine analog, Gö-6976, stimulates stress-dependent kinases in culture neonatal and adult cardiac myocytes, but not in fibroblasts. This surprising action of Gö-6976 is blocked by U-73122, which is a PLC inhibitor. Furthermore, we show that Gö-6976 alters calcium signaling in NRVMs within seconds after treatment. These findings suggest the possibility that Gö-6976 has an off-target action that serves to activate a cell surface receptor(s) to stimulate PLC-mediated calcium release (Fig. 4C).

Activation of cardiac stress signaling by Gö-6976 was initially indicated by induction of PKD phosphorylation on serine-916, which is unexpected since this is an auto-phosphorylation site, and Gö-6976 is an ATP-competitive inhibitor of PKD [16]. In this regard, it is important to stress the distinction between auto-phosphorylation of PKD and phosphorylation of downstream substrates by PKD. Rybin et al. showed that PKD1 serine-916 auto-phosphorylation does not correlate with the ability of the kinase to phosphorylate downstream substrates such as CREB [37]. Furthermore, several studies have shown that PKD-mediated phosphorylation of HDAC5 can be blocked under conditions where serine-916 phosphorylation persists [15,26,38]. Our results with catalytically inactive PKD1 (Fig. 3F) suggest that another kinase(s) is phosphorylating serine-916 in Gö-6976-treated cells. Chemical biological probes have the potential to provide key insights into the roles of specific targets in the control of various biological processes and disease pathways. Ideally, results should be confirmed with at least two independent, structurally distinct compounds that target the same protein, thereby reducing the possibility that observed effects are governed by "off-target" actions of a given small molecule. In this regard, several novel and highly selectively small molecule inhibitors of class IIa HDACs and PKD have recently been described [24,38–46]. These powerful pharmacological tools should help clarify the roles of class IIa HDAC and PKD catalytic activity in striated muscle cell differentiation and growth.

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Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at http://dx.doi.org/10.1016/j.febslet.2015.03. 017.

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