

PKD1 Inhibits AMPK α 2 through Phosphorylation of Serine 491 and Impairs Insulin Signaling in Skeletal Muscle Cells*

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Background: Diminished activity of the enzyme AMP-activated protein kinase (AMPK) is associated with impaired insulin signaling.

Results: Protein Kinase (PK)C/D1 activation inhibits AMPK α 2 via Ser⁴⁹¹ phosphorylation; PKD1 inhibition prevents this in skeletal muscle cells.

Conclusion: PKD1 is a novel upstream AMPK-kinase that phosphorylates AMPK on Ser⁴⁹¹ and regulates insulin signaling.

Significance: PKD1 inhibition may be a novel strategy for improving insulin sensitivity.

AMP-activated protein kinase (AMPK) is an energy-sensing enzyme whose activity is inhibited in settings of insulin resistance. Exposure to a high glucose concentration has recently been shown to increase phosphorylation of AMPK at Ser^{485/491} of its α 1/ α 2 subunit; however, the mechanism by which it does so is not known. Diacylglycerol (DAG), which is also increased in muscle exposed to high glucose, activates a number of signaling molecules including protein kinase (PK)C and PKD1. We sought to determine whether PKC or PKD1 is involved in inhibition of AMPK by causing Ser^{485/491} phosphorylation in skeletal muscle cells. C2C12 myotubes were treated with the PKC/D1 activator phorbol 12-myristate 13-acetate (PMA), which acts as a DAG mimetic. This caused dose- and time-dependent increases in AMPK Ser^{485/491} phosphorylation, which was associated with a ~60% decrease in AMPK α 2 activity. Expression of a phosphodeficient AMPK α 2 mutant (S491A) prevented the PMA-induced reduction in AMPK activity. Serine phosphorylation and inhibition of AMPK activity were partially prevented by the broad PKC inhibitor Gö6983 and fully prevented by the specific PKD1 inhibitor CRT0066101. Genetic knockdown of PKD1 also prevented Ser^{485/491} phosphorylation of AMPK. Inhibition of previously identified kinases that phosphorylate AMPK at this site (Akt, S6K, and ERK) did not prevent these events. PMA treatment also caused impairments in insulin-signaling through Akt, which were prevented by PKD1 inhibition. Finally, recombinant PKD1 phosphorylated AMPK α 2 at Ser⁴⁹¹ in cell-free conditions. These results identify PKD1 as a novel upstream kinase of AMPK α 2 Ser⁴⁹¹ that plays a negative role in insulin signaling in muscle cells.

Skeletal muscle is responsible for ~80% of insulin-stimulated glucose uptake in healthy humans (1). Thus, tight regulation of insulin signaling and glucose uptake in muscle cells is essential to maintaining normal blood glucose levels and overall metabolic health. This process can become dysregulated in pathological settings of insulin resistance, such as type 2 diabetes (T2D),² which currently affects over 300 million people worldwide (2). At a cellular level, prolonged exposure to excess nutrients, such as glucose, leucine, and free fatty acids (FFA), can impair the signaling processes that normally regulate glucose uptake. Though these pathways have been studied in detail, a better understanding of the pathological changes that lead to dysfunctional insulin signaling is essential for improving therapeutic strategies.

AMP-activated protein kinase (AMPK) is an energy sensing enzyme that is regarded as a master regulator of cellular metabolism (3). This serine/threonine kinase is activated when cellular energy levels are low (high AMP:ATP ratio) and its activation signals for the cell to increase catabolic (energy generating) processes, such as glucose uptake and fatty acid oxidation, and to inhibit anabolic (energy consuming) processes, such as protein and triglyceride synthesis. AMPK is activated by both its phosphorylation on Thr¹⁷² and allosteric modification of its catalytic α -subunit. Activated AMPK, in turn, mediates some of the beneficial metabolic effects of exercise and pharmacological activators, such as biguanides, and thiazolidinediones, which improve insulin sensitivity. Recent studies suggest that an inhibitory phosphorylation site at Ser^{485/491} of AMPK α 1/ α 2 subunits plays an important role in diminishing its enzyme activity (4–7). We recently found that incubation of rat extensor digitorum longus (EDL) muscle with elevated glucose levels (25 versus 5.5 mM) for 1 or 2 h both increased the phosphorylation of AMPK at Ser^{485/491} and diminished its phosphorylation at Thr¹⁷², both of which decreased its enzyme activity, as mea-

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² The abbreviations used are: T2D, type 2 diabetes; FFA, free fatty acid; diacylglycerol; PMA, phorbol 12-myristate 13-acetate; CaMK, calcium/calmodulin-dependent protein kinase; AMPK, AMP-activated protein kinase; ACC, acetyl-CoA carboxylase.

sured by the SAMS peptide assay (8). Similarly, it has been reported that when MIN6 pancreatic beta cells are switched from low glucose (3 mM) culture medium to high glucose (25 mM) for either 1 h or overnight, phosphorylation of AMPK α 1 at Ser⁴⁸⁵ is increased, and phosphorylation at Thr¹⁷² is diminished (9).

Several upstream kinases have been identified that phosphorylate AMPK at Ser^{485/491} and inhibit its activity in various tissues. For example, in heart, skeletal muscle, and liver, insulin and IGF-1 stimulate AMPK phosphorylation at this site by activating Akt (4, 7, 10, 11). Also, in the hypothalamus, Dagon *et al.* (4, 5) showed that p70S6K phosphorylates AMPK α 2 Ser⁴⁹¹ to inhibit AMPK activity and decrease food intake. Others have shown that in murine macrophage-like RAW 264.7 cells, IKK β phosphorylates AMPK at Ser⁴⁸⁵ in response to LPS treatment (12), while ERK1/2 can inhibit AMPK by this mechanism in mature dendritic cells in response to CCR7 signaling (13). Finally, protein kinase A (PKA) has been reported to phosphorylate this site in INS-1 cells in response to forskolin or GIP stimulation (14) and in human diploid fibroblasts in response to lysophosphatidic acid (15). Inhibition of AMPK through this mechanism by multiple kinases suggests a biological need to maintain tight control over AMPK enzyme activity. Whether still other kinases also phosphorylate AMPK at Ser^{485/491} to modulate its activity and biological functions remain to be determined.

Protein kinase C (PKC) is a family of serine/threonine kinases that can be activated by diacylglycerol (DAG), a phospholipid, and Ca²⁺ (depending on the isoform). Several of the 10 known PKC isoforms have long been implicated in the development of insulin resistance, but beyond inhibitory serine phosphorylation of IRS-1 (16), the mechanisms by which they do so have not been fully clarified. Protein kinase D1 (PKD1) (the mouse ortholog of human PKC μ) is a related kinase that can be activated either directly by DAG or due to phosphorylation by novel PKC isoforms (17). When activated, PKD1 undergoes transphosphorylation by novel PKCs at Ser^{744/748} of its activation loop and autophosphorylation at Ser⁹¹⁶ at its C terminus. This more recently discovered enzyme was initially classified as an atypical PKC isoform, but was later determined to be more similar structurally to the calcium/calmodulin-dependent protein kinase (CaMK) group of serine/threonine kinases with unique substrate specificity (17, 18). Pathological overactivation of PKD1 has been linked to the invasiveness of certain cancers (19) and to cardiac hypertrophy (20, 21), the latter of which is often present in patients with T2D. Whether alterations in PKD1 expression or activity affect AMPK activation is not yet known. What is clear is that the DAG mimetic phorbol 12-myristate 13-acetate (PMA), which activates PKC and PKD1, as well as many other signaling molecules, decreases AMPK activity in cardiac myocytes (22), although the mechanism by which it does so is unknown.

In the present study, we sought to determine whether PKC or PKD1 is involved in the inhibition of AMPK by phosphorylation at Ser^{485/491} in skeletal muscle cells. We used the phorbol ester PMA to determine the effects of broad PKC/PKD1 activation on AMPK phosphorylation and activity. By the use of nonspecific and specific PKC and PKD1 inhibition, we show for

the first time that PKD1 phosphorylates AMPK at Ser^{485/491}, thus diminishing AMPK activity.

Experimental Procedures

Materials—C2C12 cells were purchased from ATCC (Manassas, VA). DMEM, Penicillin-Streptomycin (P/S), fetal bovine serum (FBS), and horse serum (HS) were from Invitrogen (Grand Island, NY). D-(+)-Glucose solution, 45%, and insulin were obtained from Sigma-Aldrich. Primary antibodies for acetyl-CoA carboxylase (ACC), AMPK, phospho-AMPK α (Thr¹⁷²), phospho-AMPK α 1/ α 2 (Ser^{485/491}), phospho-PKD1 (Ser⁹¹⁶), phospho-PKD1 (Ser^{744/748}), PKD1, phospho-PKC (pan) (β II Ser⁶⁶⁰), phospho-PKC δ / θ (Ser^{643/676}), phospho-(Ser) PKC Substrate, phosphor-(Ser/Thr) PKD substrate, phospho-mTOR (Ser²⁴⁴⁸), mTOR, phospho-p70S6K (Thr³⁸⁹), phospho-ERK (Thr²⁰²/Tyr²⁰⁴), phospho-Akt (Ser⁴⁷³), and Akt antibodies, as well as secondary horseradish peroxidase (HRP)-linked antibodies were purchased from Cell Signaling Technology (Danvers, MA). Phospho-ACC (Ser⁷⁹) antibody was from Upstate/Millipore (Temecula, CA). Anti- β -actin and anti-FLAG were purchased from Sigma-Aldrich. PRKD1 siRNA and corresponding negative control scrambled siRNA were purchased from Thermo Fisher Scientific (Waltham, MA). AMPK α 1 and α 2 antibodies used for immunoprecipitation were purchased from Santa Cruz Biotechnology, Inc. SAMS peptide was purchased from Abcam (Cambridge, MA) and [γ -³²P]ATP was from Perkin-Elmer (Boston, MA).

Cell Culture—C2C12 myoblasts were cultured in normal glucose (5.5 mM) DMEM supplemented with 10% fetal bovine serum (FBS), 1% glutamine, and 1% penicillin/streptomycin (P/S). Medium was replaced every 24–48 h, and cells were passaged upon reaching 80–90% confluence. At 80–90% confluence they were differentiated into myotubes in DMEM supplemented with 2% horse serum and 1% P/S. Glucose and FBS-free DMEM, supplemented with 1% P/S and glucose at a final concentration of 5.5 mM, was used for all experimental incubations.

Transfection Studies—Cells were transfected with Lipofectamine 3000 (for FLAG-WT AMPK α 2/FLAG-S491A AMPK α 2 studies) or Lipofectamine RNAiMax (for siRNA studies) according to the manufacturer's instructions. C2C12 cells were transfected on day 3 of differentiation, and experiments were performed on day 5 of differentiation.

Cell Lysate Preparation—Cells were washed once on ice with Dulbecco's PBS, lysed in buffer containing 20 mM Tris-HCl, pH 7.5, 150 mM NaCl, 1 mM Na₂EDTA, 1% Triton, 2.5 mM sodium pyrophosphate, 1 mM β -glycerophosphate, 1 mM Na₃VO₄, 1 μ g/ml leupeptin, 1 \times phosphatase inhibitor mixture 3 (Sigma), and 1 \times protease inhibitor mixture containing 0.5 mM EDTA (Thermo Fisher Scientific), and removed from wells using a cell scraper with a polyethylene copolymer blade (Fisher Scientific). Cell debris was removed by centrifugation at 13,200 \times g for 10 min at 4 $^{\circ}$ C, and the supernatant was removed and stored at -80 $^{\circ}$ C until analysis. Protein concentration was assessed by the bicinchoninic acid method (BCA; Pierce Biotechnology, Inc., Rockford, IL).

SDS-PAGE Western Blot Analysis—Protein expression and phosphorylation were determined in 15–30 μ g of protein lysate using SDS-PAGE gel electrophoresis and immunoblotting. Fol-

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lowing transfer onto a polyvinylidene difluoride membrane, membranes were blocked in Tris-buffered saline (pH 7.5) containing 0.05% Tween-20 (*v/v*; TBST) and 5% nonfat dry milk (*w/v*) for 1 h at room temperature, followed by incubation in primary antibodies (1:1,000) at 4 °C overnight. After washing, membranes were incubated in a secondary antibody conjugated to horseradish peroxidase at a 1:5,000 dilution for 1 h at room temperature. Bands were visualized using enhanced chemiluminescence solution (ECL; Pierce Biotechnology, Inc., Rockford, IL), and densitometry was performed with Scion Image software.

AMPK Activity Assay—AMPK activity was assessed as previously described (23, 24). Briefly, AMPK α 1 or α 2 was immunoprecipitated from 500 μ g of protein from cell lysates by incubation at 4 °C overnight on a roller mixer using AMPK α 1 or α 2-specific antibodies (1:80) and protein A/G-agarose beads (1:10; Santa Cruz Biotechnology, Inc). Following several washes, activity was measured in the presence of 200 μ M AMP and 80 μ M [γ -³²P]ATP (2 μ Ci) using 200 μ M SAMS peptide (Abcam) as a substrate. Label incorporation into the SAMS peptide was quantified using a LabLogic (Brandon, FL) scintillation counter.

Cell-free in Vitro Phosphorylation Assay—Recombinant proteins were purchased from EMD Millipore (Billerica, MA) (AMPK) and Enzo Life Sciences (Farmington, NY) (PKD1 and Akt). Recombinant α 2AMPK/ β 1/ γ 1 complex was incubated with recombinant Akt or PKD1 in 50 mM Na-HEPES, 5 mM MgCl₂, 500 μ M ATP, 1 mM DTT for 30 min at 30 °C.

Statistical Analysis—Results are reported as means \pm S.E. of the mean (S.E.). Statistical significance was determined by two-tailed unpaired Student's *t* tests or ANOVA with Tukey's post-hoc test. A level of *p* < 0.05 was considered statistically significant.

Results

The PKC Activator PMA Dose- and Time-dependently Stimulates AMPK Ser^{485/491} Phosphorylation in C2C12 Myotubes—Treatment with the PKC activator phorbol 12-myristate 13-acetate (PMA) has previously been shown to decrease AMPK activity in cardiac myocytes, as measured by the SAMS peptide assay (22). We sought to determine whether PMA had the same effect in C2C12 myotubes, and if so, whether it is mediated by an inhibitory phosphorylation on AMPK α 1/ α 2 subunit at Ser^{485/491}. All doses of PMA (10–100 nM) significantly increased serine phosphorylation of AMPK (Fig. 1, *A* and *B*), as did PMA treatment for 30 min to 3 h (50 nM) (Fig. 1, *C* and *D*). Increases in overall PKC activity, as measured by serine phosphorylation of PKC substrates (data not shown), mirrored the increases in phosphorylation of AMPK Ser^{485/491}. Phosphorylation of PKD at Ser⁹¹⁶ and Ser^{744/748} of its activation loop were also significantly increased by incubation with PMA at doses of 25 nM or higher for 30 min to 3 h (Fig. 1, *A–D*). Since phosphorylation of these two sites changed similarly in response to all treatments, only the data for Ser⁹¹⁶ phosphorylation is shown for subsequent experiments. Notably, no significant changes in phosphorylation of AMPK Thr¹⁷² (Fig. 1, *A* and *C*) or its downstream substrate ACC Ser⁷⁹ (data not shown) were observed. Phosphorylation of conventional PKC isoforms, as measured

by p-PKC (pan) (β II Ser⁶⁶⁰), was increased in a dose-response manner; however, activation of the atypical PKC ζ was not increased by PMA treatment (data not shown).

The Non-selective PKC Inhibitor Gö6983 Prevents PMA-induced Phosphorylation of AMPK Ser^{485/491} in the Myotubes—To determine whether the increase in phosphorylation of AMPK at Ser^{485/491} was attributable to PKC activation, we used the non-selective PKC inhibitor Gö6983. Treatment of C2C12 myotubes with Gö6983 (5 μ M) for 1 h prior to PMA treatment (50 nM, 30 min) significantly attenuated phosphorylation of AMPK at Ser^{485/491} (Fig. 2, *A* and *B*). Phosphorylation of AMPK Thr¹⁷² (Fig. 2*A*) was not affected by PMA treatment, but it was increased by the inhibitor. PKC activity, assessed by serine phosphorylation of PKC substrates, was decreased by the inhibitor (Fig. 2*A*), as was phosphorylation of conventional PKC isoforms, when evaluated by phospho-PKC β pan, and PKD at Ser⁹¹⁶ (Fig. 2, *A* and *B*), but not PKC δ/θ (data not shown).

The Selective PKD1 Inhibitor CRT0066101 Prevents PMA-induced Phosphorylation of AMPK Ser^{485/491}—Since phosphorylation of the activation loop of PKD at Ser^{744/748} and at Ser⁹¹⁶ (Fig. 2, *A* and *B*) corresponded with changes in phosphorylation of AMPK at Ser^{485/491}, we investigated whether specific inhibition of PKD could attenuate PMA-induced serine phosphorylation of AMPK. Pre-treatment for 1 h with the specific PKD inhibitor CRT0066101 (10 μ M) prevented phosphorylation of AMPK Ser^{485/491} in C2C12 myotubes (Fig. 2, *C* and *D*). Phosphorylation of AMPK at Thr¹⁷² was unchanged by treatment with this inhibitor (Fig. 2*C*). As expected, PMA effects on PKD Ser⁹¹⁶ phosphorylation were ablated by CRT0066101 treatment, and probing with a phospho-(Ser/Thr) PKD substrate antibody revealed an overall increase in number of bands and band intensity with PMA treatment, which was prevented by pretreatment with CRT0066101 (Fig. 2*C*). Bands detected by this antibody at 110 kDa and 62 kDa may represent PKD (which is autophosphorylated at Ser⁹¹⁶) and AMPK, respectively.

PMA Treatment Inhibits AMPK Activity, which Is Attenuated by Pretreatment with Gö6983 or CRT0066101—Recent studies have shown that phosphorylation of AMPK Ser^{485/491} can inhibit its activity, even in the absence of diminished Thr¹⁷² phosphorylation, which is often used as a surrogate readout for enzyme activity (4, 5). We sought to determine whether the changes in phosphorylation of AMPK Ser^{485/491} caused by PMA and the PKC/D inhibitors Gö6983 and CRT0066101 corresponded to changes in AMPK activity. Using the SAMS peptide assay described in the methods section, we found that PMA treatment significantly decreased AMPK α 2 activity in C2C12 myotubes (Fig. 3, *A* and *C*). Furthermore, pretreatment with the non-selective PKC inhibitor Gö6983 significantly attenuated the reduction in enzyme activity (Fig. 3*A*), while the specific PKD inhibitor CRT0066101 completely prevented the PMA-induced decrease in AMPK activity (Fig. 3*C*). Despite the lack of changes in phosphorylation of AMPK Thr¹⁷² or ACC Ser⁷⁹, we found a significant inverse correlation between AMPK α 2 activity and phosphorylation at Ser^{485/491} (Fig. 3, *B* and *D*).

Phosphorylation of AMPK α 2 on Ser⁴⁹¹ Is Necessary for PMA-induced Inhibition of AMPK Activity—To determine if serine phosphorylation is required for the PMA-induced reduction in AMPK activity, C2C12 cells were transfected with WT

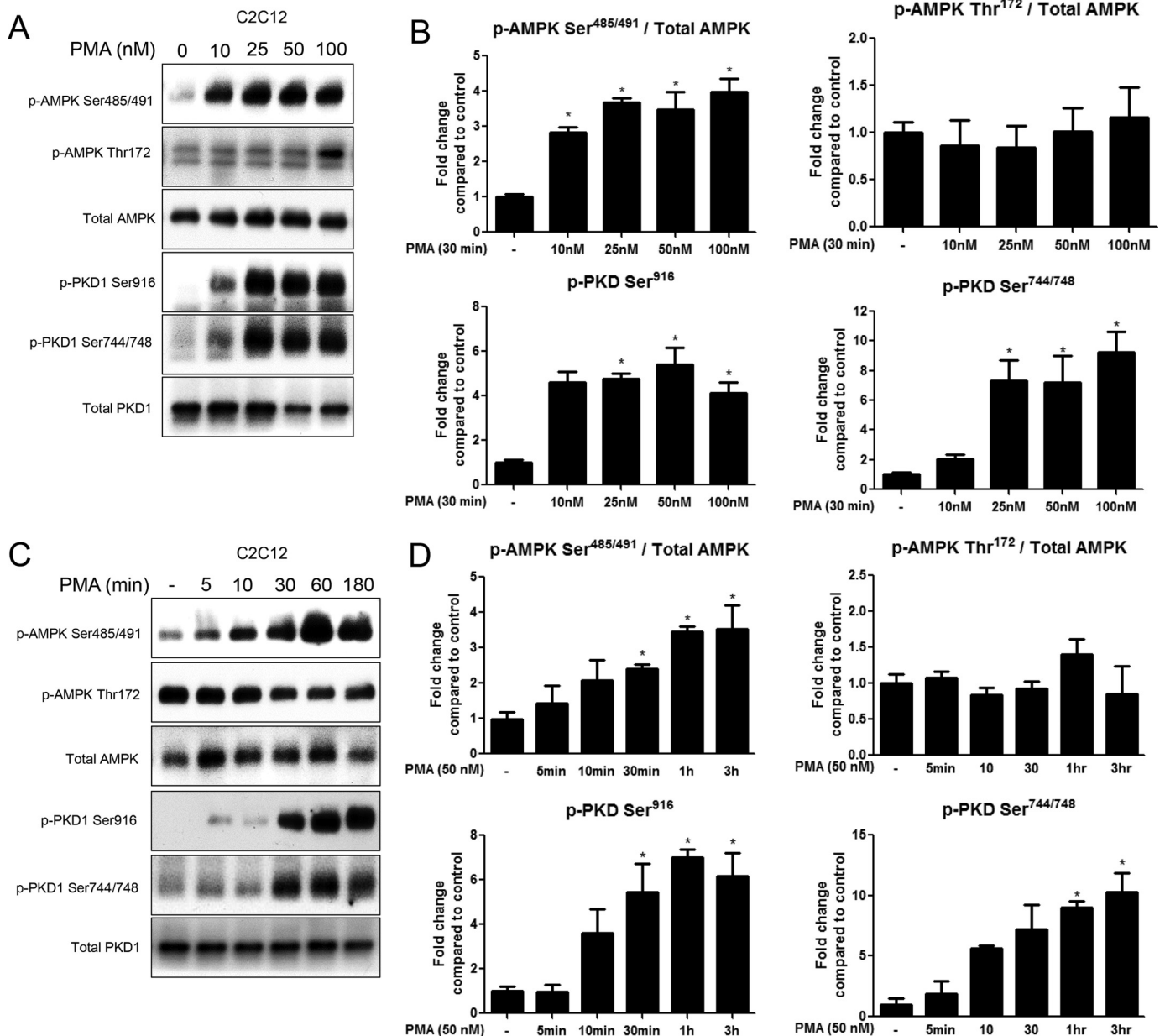


FIGURE 1. PMA treatment time- and dose-dependently stimulates phosphorylation of AMPK at Ser^{485/491} in C2C12 myotubes. Myotubes were treated with 10–100 nM PMA for 30 min (A, B) or 50 nM PMA for 5–180 min (C, D). Cells were lysed and protein expression and phosphorylation were analyzed by Western blot. Representative Western blots (A, C) and their densitometric analyses (B, D) are shown. Results are means \pm S.E. ($n = 3$ –6 per treatment). All experiments were performed in triplicate. *, $p < 0.05$ compared with control.

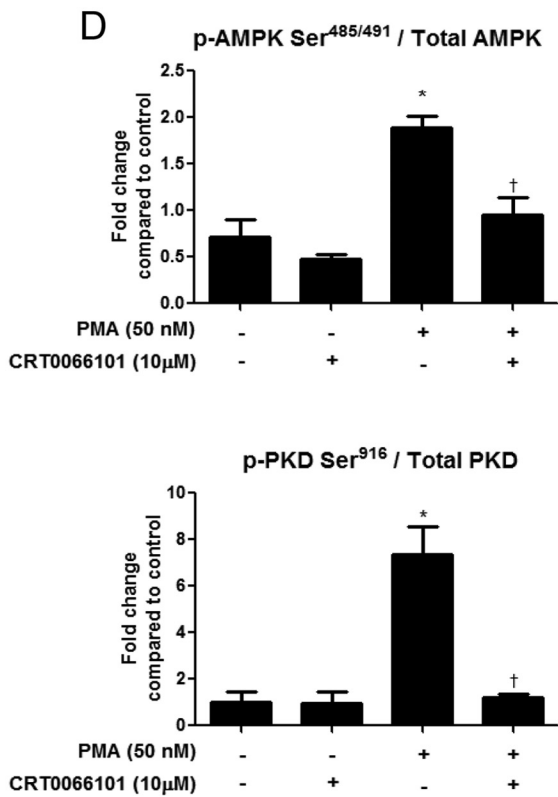
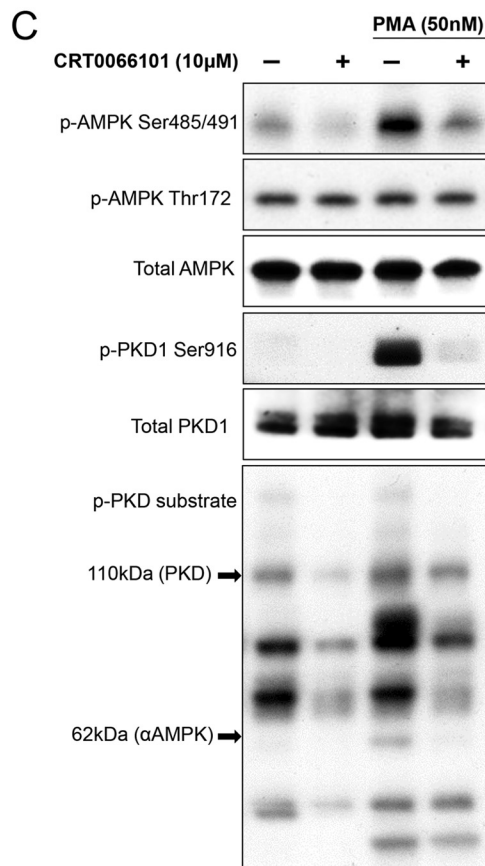
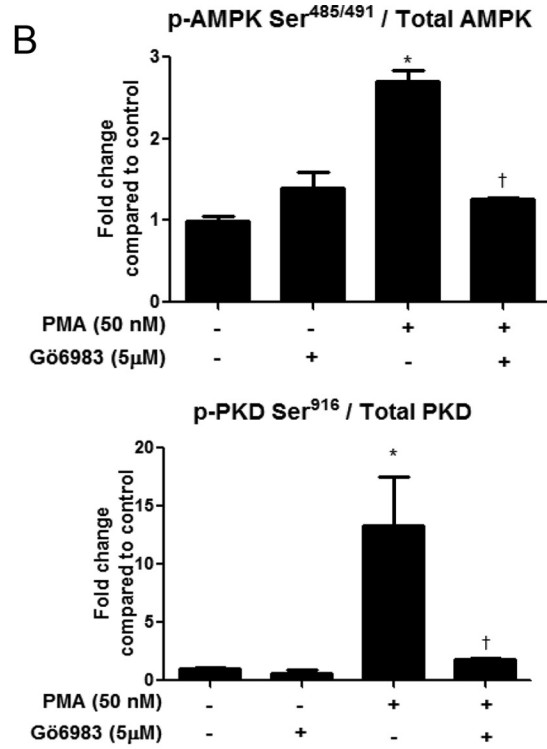
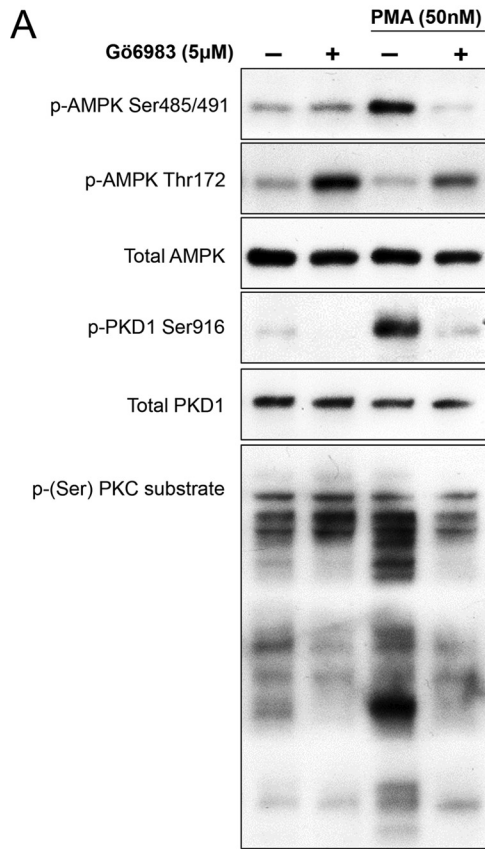
AMPK α 2 or phosphodeficient S491A AMPK α 2. As expected, phosphorylation of AMPK at Ser⁴⁹¹ under both basal and PMA-treated conditions was prevented in the S491A AMPK α 2-expressing cells (Fig. 3E). While PMA significantly decreased AMPK α 2 activity, as measured by the SAMS peptide assay, in cells transfected with WT AMPK α 2, no effect of PMA treatment was observed in the S491A AMPK α 2-expressing cells. These data suggest that Ser⁴⁹¹ phosphorylation of AMPK α 2 is required for the PMA-induced inhibition of AMPK activity.

Knockdown of PKD1 Prevents PMA-induced Phosphorylation of AMPK Ser^{485/491}—To confirm the results obtained using pharmacological inhibitors which suggested that PKD1 is involved in PMA-induced AMPK Ser^{485/491} phosphorylation (Fig.

2), we used siRNA to reduce PKD1 mRNA, and hence protein, levels. As expected, transfection of C2C12 cells with PKD1 (gene name PRKD1) caused a reduction in phospho-PKD1 (Fig. 4, A and B). Consistent with data shown in Fig. 2, PKD1 knockdown prevented the PMA-induced increase in AMPK Ser^{485/491} phosphorylation. These data offer further evidence that PKD1 is responsible for PMA-induced inhibition of AMPK by Ser^{485/491} phosphorylation.

Inhibition of Akt, P70S6K, or ERK Does Not Prevent PMA-induced Phosphorylation of AMPK Ser^{485/491}—As previously mentioned, the kinases Akt, P70S6K, and ERK1/2 have been shown to phosphorylate AMPK at Ser^{485/491} in response to various stimuli in other cell types. To rule out involvement of these kinases in PMA-induced phosphorylation of AMPK Ser^{485/491},

PKD1 Inhibits AMPK α 2 Activity via Phosphorylation on Ser⁴⁹¹



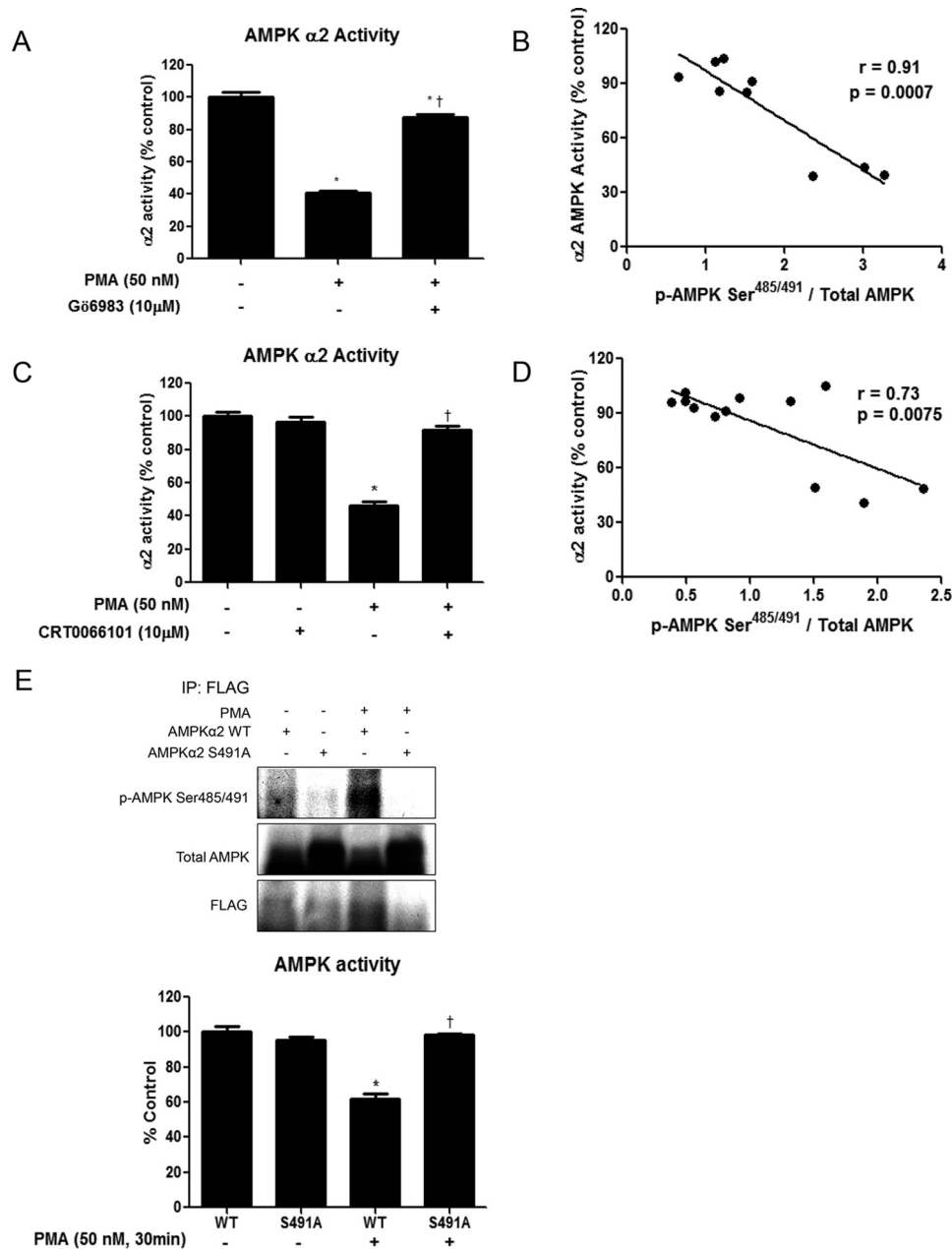


FIGURE 3. PMA treatment diminishes AMPK activity in C2C12 myotubes, while PKC and PKD inhibition or expression of S491A AMPK α 2 prevent this decrease. C2C12 myotubes were treated with the nonspecific PKC inhibitor Gö6983 (5 μ M) (A, B) or the specific PKD inhibitor CRT0066101 (10 μ M) (C, D) for 1 h prior to PMA treatment (50 nM, 30 min). AMPK α 2 activity was measured by the SAMS peptide assay. B and D show the correlation between AMPK α 2 activity and Ser^{485/491} phosphorylation. E, C2C12 cells were transfected with FLAG-tagged WT AMPK α 2 or S491A AMPK α 2. 48 h later, cells were treated with PMA (50 nM, 30 min) and then lysed and immunoprecipitated using an anti-FLAG antibody. Protein phosphorylation and expression were analyzed by Western blot. AMPK activity was measured using the SAMS peptide assay. Results are means \pm S.E. ($n = 3-6$ per treatment). *, $p < 0.05$ compared with control; †, $p < 0.05$ compared with PMA treatment.

we used inhibitors of each of these kinases. Pretreatment of C2C12 myotubes with the PI3K/Akt inhibitor wortmannin (100 nM) did not affect PMA-stimulated serine phosphorylation of AMPK (Fig. 5A). As expected, PMA treatment did not increase Akt phosphorylation, but p-Akt Ser⁴⁷³ phosphorylation was diminished by wortmannin treatment. Next, we used

the mTOR inhibitor rapamycin to test whether P70S6K was involved. Although P70S6K Thr³⁸⁹ phosphorylation was increased by PMA treatment, rapamycin (100 nM) treatment did not affect PMA-induced phosphorylation of AMPK Ser^{485/491} (Fig. 5B). Similarly, pretreatment with the ERK inhibitor U0126 (5 μ M) had no effect on AMPK serine phosphoryla-

FIGURE 2. PKD inhibition prevents PMA-induced phosphorylation of AMPK Ser^{485/491} in C2C12 myotubes. Myotubes were treated with the nonspecific PKC inhibitor Gö6983 (5 μ M) (A, B) or the specific PKD inhibitor CRT0066101 (10 μ M) (C, D) for 1 h prior to PMA treatment (50 nM, 30 min). Cells were lysed and protein expression and phosphorylation were analyzed by Western blot. Representative Western blots (A, C) and their densitometric analyses (B, D) are shown. Results are means \pm S.E. ($n = 3-6$ per treatment). All experiments were performed in triplicate. *, $p < 0.05$ compared with control; †, $p < 0.05$ compared with PMA treatment.

PKD1 Inhibits AMPK α 2 Activity via Phosphorylation on Ser⁴⁹¹

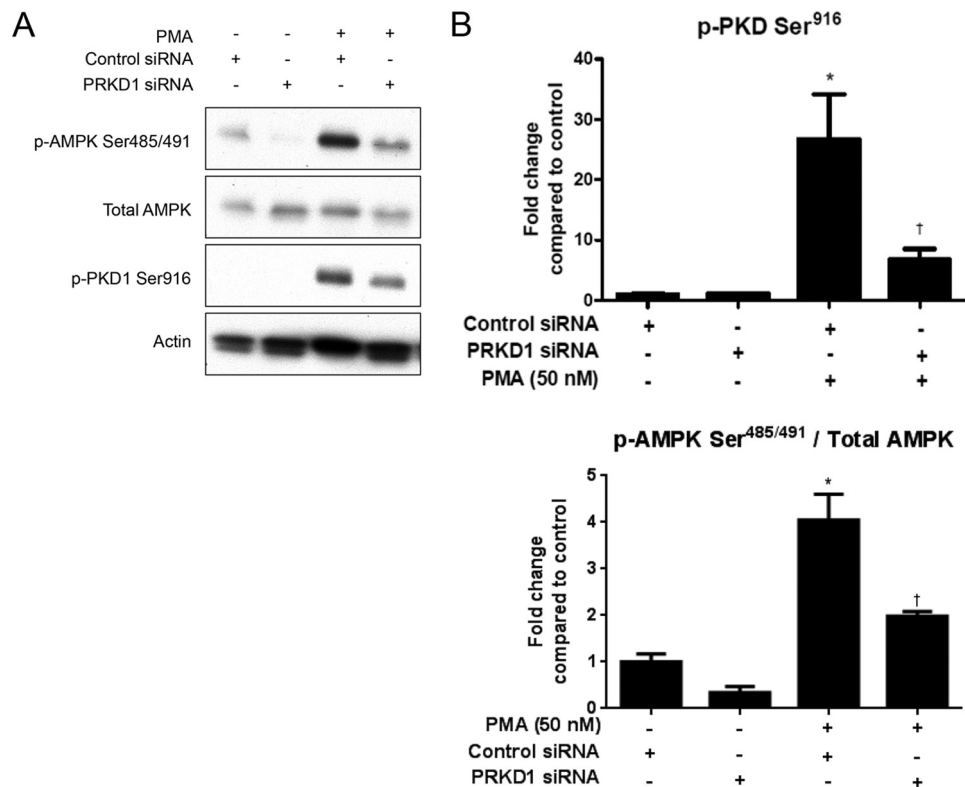


FIGURE 4. PKD1 knockdown prevents PMA-induced phosphorylation of AMPK Ser^{485/491} in C2C12 myotubes. C2C12 myotubes were transfected with scrambled or PKD1 siRNA. 48 h later, they were treated with PMA (50 nM, 30 min). Cells were lysed, and protein expression and phosphorylation were analyzed by Western blot. Representative Western blots (A) and their densitometric analyses (B) are shown. Results are means \pm S.E. ($n = 3$ –6 per treatment). All experiments were performed in triplicate. *, $p < 0.05$ compared with control; †, $p < 0.05$ compared with PMA treatment.

tion (Fig. 5C), despite inhibiting ERK phosphorylation, which can occur downstream of PKD.

PMA Treatment Inhibits Insulin Signaling through Akt, which Is Attenuated by Pretreatment with Gö6983 or CRT0066101, or by PKD1 Knockdown—To test whether the PMA-induced changes were associated with impaired insulin signaling, we measured insulin-stimulated Akt phosphorylation. PMA treatment (50 nM) for 30 min diminished insulin-stimulated Akt Ser⁴⁷³ phosphorylation (Fig. 6, A and B). Pretreatment with Gö6983 or CRT0066101 restored insulin signaling through Akt. Since both insulin and PMA independently stimulate AMPK Ser^{485/491} phosphorylation through Akt and PKD, respectively, all combinations of insulin and PMA with and without inhibitors significantly increased serine phosphorylation of AMPK (Fig. 6, A and B). Similarly, siRNA mediated knockdown of PKD1 also restored insulin signaling through Akt (Fig. 6, C and D). PKD1 knockdown, however, did not prevent PMA-induced serine phosphorylation of IRS-1, which is known to impair insulin signaling. Thus, this suggests that PKD1 activation impairs insulin signaling through a novel mechanism, independent of IRS-1 phosphorylation.

Expression of S491A AMPK α 2 Prevents PMA-induced Impairments in Insulin-signaling through Akt—To evaluate whether AMPK was involved in the diminished insulin-stimulated Akt phosphorylation following PMA treatment, we transfected cells with phosphodeficient S491A AMPK α 2. While PMA treatment significantly reduced insulin-stimulated Akt Ser⁴⁷³ phosphorylation in cells expressing WT AMPK α 2, this effect was not seen in cells expressing the phosphodeficient

AMPK α 2 (Fig. 7). These data suggest that AMPK Ser⁴⁹¹ is required for PMA to impair insulin signaling through Akt.

Recombinant PKD1 Phosphorylates AMPK α 2 at Ser⁴⁹¹ in Cell-free Conditions—To determine if PKD1 directly phosphorylates AMPK or if an intermediate signaling molecule is necessary, we tested whether recombinant PKD1 phosphorylates AMPK in a cell-free system. Compared with AMPK α 2 alone, co-incubation of recombinant PKD1 with AMPK α 2 caused phosphorylation of AMPK α 2 at Ser⁴⁹¹ (Fig. 8A). Incubation of Akt, used as a positive control, with AMPK also caused an obvious increase in Ser⁴⁹¹ phosphorylation. These data suggest that PKD1 phosphorylates AMPK directly. To evaluate whether AMPK α 2 Ser⁴⁹¹ is a good substrate for PKD1, we lined up the PKD1 substrate consensus sequence with the sequence of residues preceding Ser⁴⁹¹ on AMPK α 2. Although AMPK α 2 has a proline rather than PKD1 preferred leucine at the -5 position, the arginine at the -3 position does line up (Fig. 8B). This is similar to Akt, which prefers arginine at both the -5 and -3 positions.

Discussion

A better understanding of how AMPK is regulated at a molecular level is important for improving drug-targeting strategies for T2D. Phosphorylation of AMPK at Thr¹⁷² of the activation loop is generally held to be essential for the activation of AMPK and is often used as a readout for its activity. However, our data add to a growing number of studies showing that phosphorylation of AMPK at Ser^{485/491} can inhibit AMPK activity, even in the absence of changes in p-AMPK Thr¹⁷² (4, 5). These

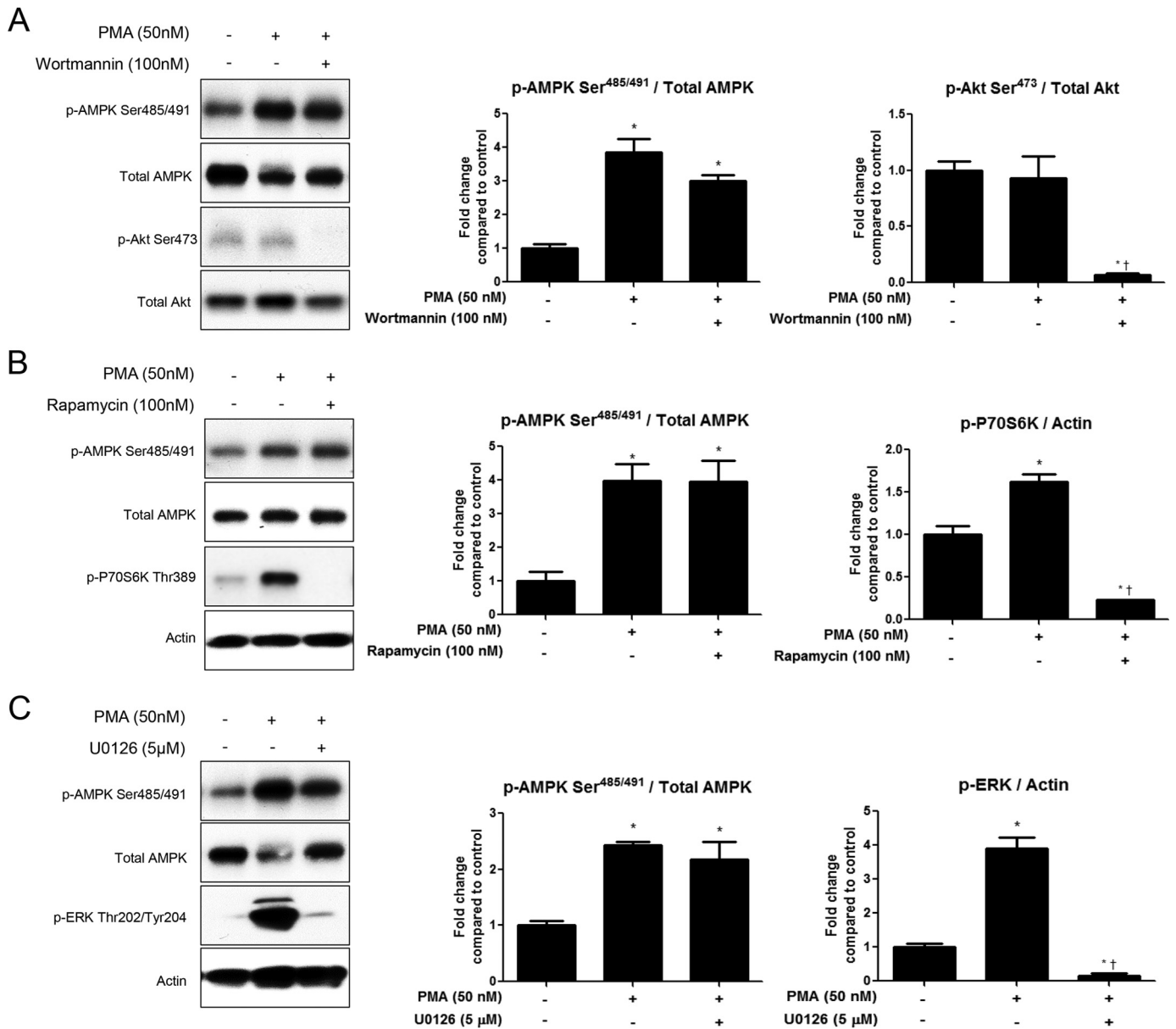


FIGURE 5. Inhibition of Akt, S6K, or ERK does not prevent PMA-induced phosphorylation of AMPK Ser^{485/491}. Myotubes were treated with the PI3K/Akt inhibitor wortmannin (100 nM) (A), the mTOR/P70S6K inhibitor rapamycin (100 nM) (B), or the ERK inhibitor U0126 (5 μ M) (C) for 2 h prior to PMA treatment (50 nM, 30 min). Cells were lysed, and protein expression and phosphorylation were analyzed by Western blot. Representative Western blots and their densitometric analyses are shown. Results are means \pm S.E. ($n = 3-6$ per treatment). All experiments were performed in triplicate. *, $p < 0.05$ compared with control; †, $p < 0.05$ compared with PMA treatment.

data suggest that phosphorylation of AMPK at Ser^{485/491} can inhibit AMPK activity independent of changes in p-AMPK Thr¹⁷² and that inhibition of p-AMPK Thr¹⁷² is not required for Ser^{485/491} phosphorylation to induce inhibition of AMPK activity. We also show, for the first time, that PKD1 activation can stimulate phosphorylation of AMPK Ser^{485/491} in skeletal muscle cells, resulting in impaired insulin signaling. In addition to identifying a novel kinase that can inhibit AMPK via Ser^{485/491} phosphorylation, these studies present a possible mechanism by which AMPK may be inhibited by excess nutrients, since DAG (which activates PKD1) is known to be increased by high glucose and lipids.

In 2012, Tsuchiya *et al.* (22) demonstrated that 60 min of PMA treatment (1 μ M) inhibits AMPK activity by \sim 33% in

cardiac myocytes, as measured by the SAMS peptide assay. They found that this inhibition could be prevented by the non-specific PKC inhibitor bisindolylmaleimide I (BIM I) and by the ERK inhibitor U0126, which led them to conclude that PMA inhibits AMPK by a mechanism involving the PKC and ERK pathways. There was no apparent change in energy state (AMP/ATP ratio) accompanying this decrease in AMPK activity. To our knowledge, this was the first report of PMA treatment causing inhibition of AMPK; however, AMPK phosphorylation at Thr¹⁷² or Ser^{485/491} was not measured. In concert with these results, we found that PMA treatment significantly diminished AMPK activity in C2C12 myotubes (Fig. 3A, 3C). However, we found that a dose of only 50 nM for 30 min resulted in a 50–60% decrease in AMPK activity, nearly twice as much as was seen in

PKD1 Inhibits AMPK α 2 Activity via Phosphorylation on Ser⁴⁹¹

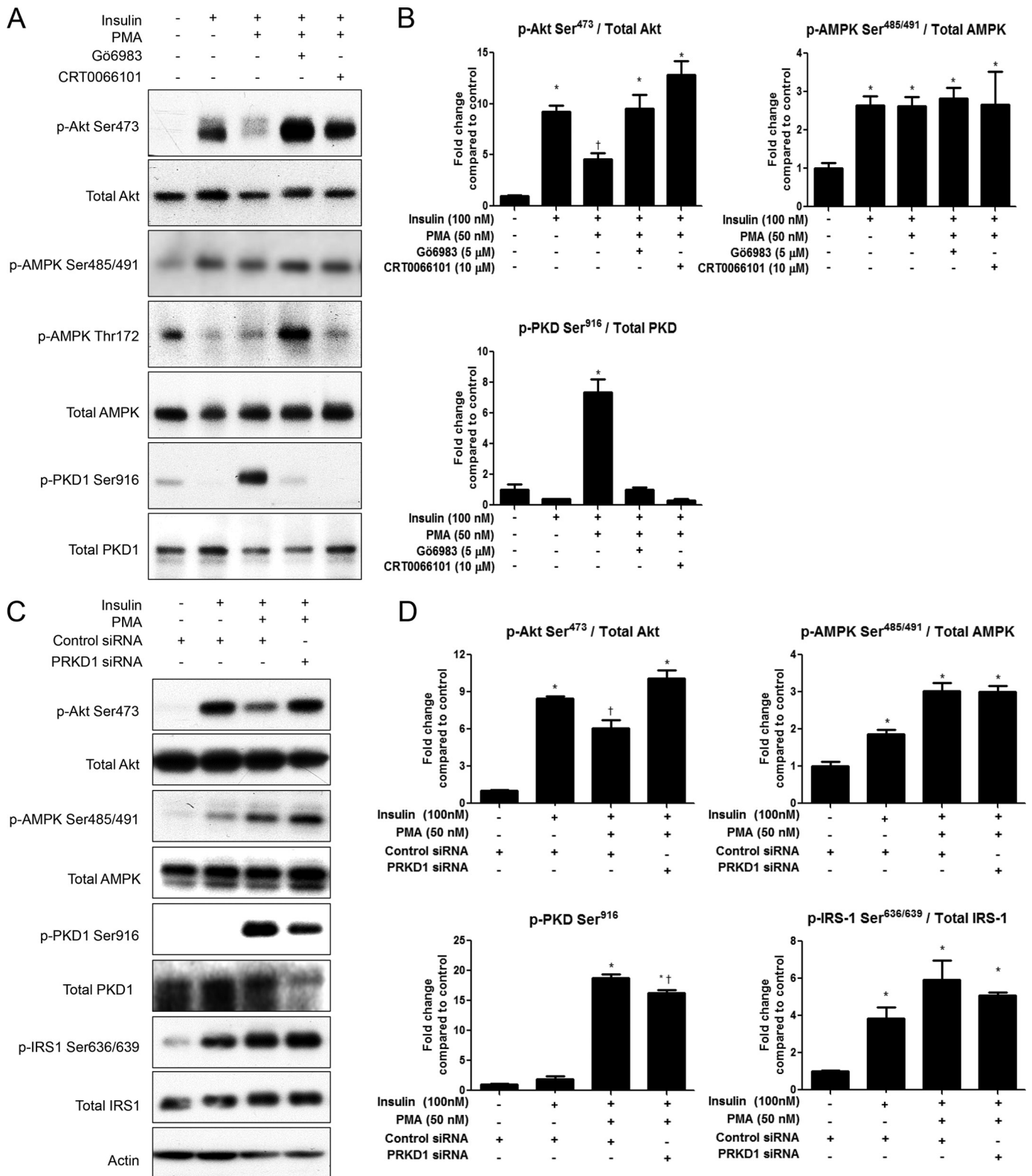


FIGURE 6. PMA stimulates PKD1 to impair insulin signaling. A and B, C2C12 myotubes were treated with or without Gö6983 (5 μ M) or CRT0066101 (10 μ M) for 1 h prior to PMA treatment (50 nM, 30 min), followed by insulin stimulation (100 nM, 15 min). C and D, C2C12 myotubes were transfected with scrambled or PKD1 siRNA. 48 h later, they were treated with PMA (50 nM, 30 min), followed by insulin stimulation (100 nM, 15 min). Cells were lysed, and Akt, AMPK, PKD1, and IRS-1 phosphorylation were analyzed by Western blot. Representative blots (A, C) and quantifications (B, D) are shown. *, $p < 0.05$ compared with control; †, $p < 0.05$ compared with insulin treatment.

PKD1 Inhibits AMPK α 2 Activity via Phosphorylation on Ser⁴⁹¹

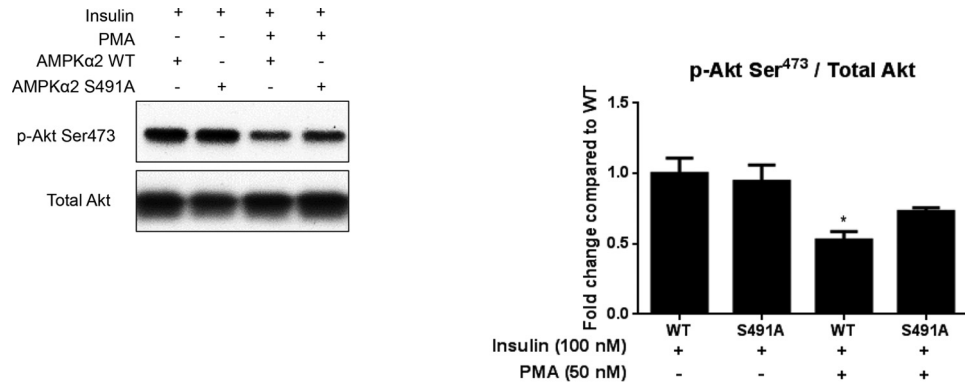


FIGURE 7. **Expression of S491A AMPK α 2 prevents the PMA-induced impairment in insulin signaling through Akt.** C2C12 cells were transfected with WT AMPK α 2 or S491A AMPK α 2. 48 h later, cells were treated with PMA (50 nM, 30 min) followed by insulin stimulation (100 nM, 15 min). Cells were then lysed, and Akt phosphorylation at Ser⁴⁷³ was analyzed by Western blot. A representative blot and quantification are shown. Results are means \pm S.E. ($n = 3-6$ per treatment). *, $p < 0.05$ compared with WT + insulin.

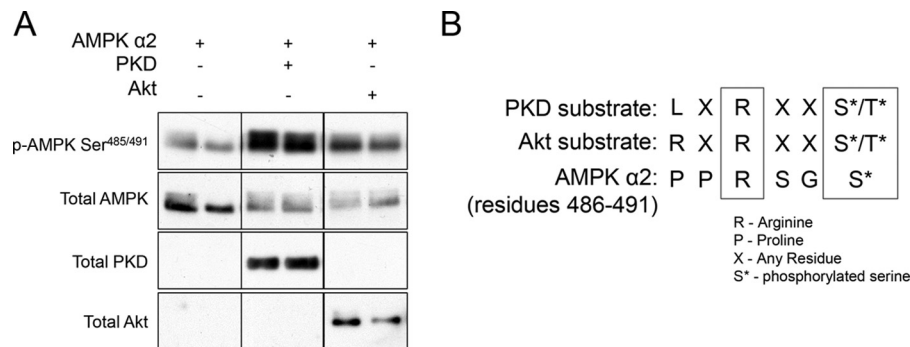


FIGURE 8. **Recombinant PKD1 phosphorylates AMPK at Ser^{485/491} in a cell-free assay.** Recombinant AMPK α 2/ β 1/ γ 1 was incubated with recombinant PKD1 or recombinant Akt as described under "Experimental Procedures." Phosphorylation of AMPK at Ser⁴⁹¹ was evaluated by Western blot. Representative image shown is from a single gel/exposure. A, consensus substrate motifs of PKD and Akt were lined up with residues 486-491 of AMPK α 2 (B). $n = 3$ per group. All experiments were performed in triplicate.

the cardiac myocytes. In addition, our results indicate that ERK was not involved in the phosphorylation of AMPK at Ser^{485/491} induced by PMA. These discrepancies could be due to differences in cell type or methodologies.

Several kinases have been reported to phosphorylate AMPK at Ser^{485/491}. We (4) and others (7) have shown that Akt phosphorylates and inhibits AMPK at this site in muscle, liver, and heart in response to insulin. In the hypothalamus, the mTOR/p70S6 kinase pathway has been reported to cause this inhibitory event in response to leptin (5), an effect shown to be essential for mediating the anorectic effects of leptin. In murine macrophage-like RAW 264.7 cells, Park *et al.* (12) showed that IKK β phosphorylates AMPK at Ser⁴⁸⁵ in response to LPS treatment in mature dendritic cells. ERK1/2 was recently shown to phosphorylate AMPK on this serine residue (13), an effect that may be an important part of the mechanism by which CCR7 signaling promotes cell survival. Furthermore, protein kinase A (PKA) has been reported to act as an upstream kinase for AMPK Ser^{485/491} in INS-1 cells in response to forskolin or GIP stimulation (14) and in human diploid fibroblasts in response to lysophosphatidic acid (15). We ruled out Akt, S6K, and ERK as upstream kinases responsible for AMPK Ser^{485/491} phosphorylation in response to PMA treatment (Fig. 5); however, we have identified PKD1 as a novel enzyme to be added to the growing list of kinases which inhibit AMPK via Ser^{485/491} phosphorylation. Furthermore, we found a significant inverse correlation

between AMPK activity and phosphorylation at Ser^{485/491} under these conditions, suggesting that PKD1 is responsible for the serine phosphorylation and inhibition of AMPK in this setting (Fig. 3, B and D). The fact that expression of a phosphodeficient S491A AMPK α 2 mutant prevented the PMA-induced reduction in AMPK α 2 activity confirms that phosphorylation of AMPK α 2 on Ser⁴⁹¹ is required for PMA-induced inhibition of AMPK kinase activity.

Many of the aforementioned kinases that can mediate AMPK inhibition through Ser^{485/491} phosphorylation are associated with anabolic or pro-hypertrophic signaling pathways (Akt, mTOR/P70S6K). This is not surprising, since AMPK is generally activated under low energy conditions that favor catabolic processes, and inhibited under conditions of energy overload that favor anabolic processes. Consistent with this, PKD1 has recently been identified as having pro-hypertrophic functions. For example, Harrison *et al.* (21) reported that transgenic mice expressing a cardiac-specific, constitutively active PKD1 mutant develop cardiac hypertrophy, followed by ventricular chamber dilation, wall thinning, and contractile dysfunction. They also showed that PKD1 expression and activity are significantly increased in the myocardium of spontaneously hypertensive rats with heart failure, an effect that is further exaggerated by thoracic aortic banding. Furthermore, PKD1 activation was found to be elevated in hearts of humans with idiopathic dilated cardiomyopathy (20). While these results suggest that

PKD1 activation may be sufficient to induce pathological cardiac remodeling, the role of PKD1 in skeletal muscle is less clear. McGee *et al.* (25) reported that loss of AMPK in muscle resulted in no change in HDAC5 phosphorylation during exercise, but a ~33% compensatory increase in PKD activation, suggesting potential redundancies in the functions of these two proteins in response to exercise. Although they did not investigate the effects of PKD loss or activation on AMPK activation, these data, together with our results, suggest that PKD and AMPK may negatively regulate each other in skeletal muscle.

Further evidence supporting potential redundancy of functions between AMPK and PKD1, at least in regards to HDAC5 phosphorylation in response to exercise, is the fact that mice overexpressing constitutively active PKD1 in skeletal muscle show resistance to fatigue in response to repetitive contraction and a shift to Type I and Type IIa muscle fibers (26), while expression of a dominant negative kinase dead PKD1 in skeletal muscle results in impaired running performance and prevents fiber type switching (27). Similarly, it has long been known that AMPK mediates many adaptations of muscle to exercise (28), and loss of AMPK in muscle causes impairments in running capacity (29).

Our data showed that PKD1 activation impairs insulin-signaling through Akt in C2C12 myotubes, an effect prevented both by pharmacological inhibition and by genetic knockdown of PKD1 (Fig. 6). PKC activation by PMA is known to impair insulin signaling by causing serine phosphorylation of IRS-1. Intriguingly, PKD1 knockdown did not reduce PMA-induced serine phosphorylation of IRS-1, despite restoring insulin-stimulated Akt phosphorylation. These data suggest that PKD1 activation impairs insulin-signaling through an IRS-1-independent mechanism, although the exact mechanism remains to be determined. Additionally, we have shown that AMPK α 2 Ser⁴⁹¹ is required for the observed impairment in insulin signaling, though further studies are required to identify the exact mechanism (Fig. 7). Although AMPK α 2 is the predominant isoform in skeletal muscle, the presence of AMPK α 1 Ser⁴⁸⁵ could explain why Akt phosphorylation is not fully restored to control levels. The present study is the first report of PKD1 activation causing impaired insulin-signaling, and the mechanism by which it does so requires further investigation, though we have shown that it involves AMPK.

In summary, we have reported a novel inhibitory relationship between PKD1 and AMPK in skeletal muscle cells whereby PKD1 inhibits AMPK directly by phosphorylating it at Ser⁴⁹¹ of the α 2 subunit. Since PKD1 may be activated directly by DAG (or the DAG mimetic PMA) or by novel PKC isoforms, we have not ruled out the possibility that novel PKC isoforms may also be involved upstream of PKD1 in these processes. Similarly, though we have shown that PKD1 activation is necessary for the inhibition of AMPK and that AMPK α 2 Ser⁴⁹¹ is required for the impairment of insulin signaling through Akt, further studies are required to further elucidate the pathway responsible for this impairment. Our data suggest that PKD1 inhibition may be a novel strategy for preventing AMPK inhibition and impaired insulin sensitivity in muscle; however, based on the growing list of PKD1 functions in a variety of tissues being reported in the

literature, this strategy would have to be approached with caution, and likely in a highly controlled, tissue-specific manner.

Author Contributions—K. A. C. and R. J. V. designed, performed, and analyzed the experiments, and K. A. C. wrote the manuscript. K. A. and B. S. S. provided technical assistance in performing experiments and preparation of figures. N. B. R. and A. K. S. helped with experimental design and editing of the manuscript. Y. D. provided reagents and technical guidance for data shown in Figs. 3 and 6. BBK edited the manuscript. All authors reviewed the results and approved the final version of the manuscript.

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