

Published in final edited form as:

J Cell Physiol. 2009 May ; 219(2): 402. doi:10.1002/jcp.21684.

Dystrophin Glycoprotein Complex-associated G $\beta\gamma$ Subunits Activate Phosphatidylinositol-3-Kinase/Akt signaling in Skeletal Muscle in a Laminin-dependent Manner

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Abstract

Previously, we showed that laminin-binding to the dystrophin glycoprotein complex (DGC) of skeletal muscle causes a heterotrimeric G-protein, (G $\alpha\beta\gamma$) to bind, changing the activation state of the G α subunit. Others have shown that laminin-binding to the DGC also leads to Akt activation. G $\beta\gamma$, released when G α is activated, is known to bind phosphatidylinositol 3-kinase (PI3K), which activates Akt in other cells. Here, we investigate whether muscle Akt activation results from G $\beta\gamma$, using immunoprecipitation and immunoblotting, and purified G $\beta\gamma$. In the presence of laminin, PI3K-binding to the DGC increases and Akt becomes phosphorylated and activated (pAkt), and glycogen synthase kinase is phosphorylated. Antibodies, which specifically block laminin-binding to α -dystroglycan, prevent PI3K-binding to the DGC. Purified bovine brain G $\beta\gamma$ also caused PI3K and Akt activation. These results show that DGC-G $\beta\gamma$ is binding PI3K and activating pAkt in a laminin-dependent manner. *Mdx* mice, which have greatly diminished amounts of DGC proteins, display elevated pAkt signaling and increased expression of integrin β 1 compared to normal muscle. This integrin binds laminin, G $\beta\gamma$, and PI3K. Collectively, these suggest that PI3K is an important target for the G $\beta\gamma$, which normally binds to DGC syntrophin, and activates PI3K/Akt signaling. Disruption of the DGC in *mdx* mouse is causing dis-regulation of the laminin-DGC-G $\beta\gamma$ -PI3K-Akt signaling and is likely to be important to the pathogenesis of muscular dystrophy. Up-regulating integrin β 1 expression and activating the PI3K/Akt pathway in muscular dystrophy may partially compensate for the loss of the DGC. The results suggest new therapeutic approaches to muscle disease.

Keywords

Duchenne muscular dystrophy (DMD); dystrophin–glycoprotein complex (DGC); phosphoinositide-3-kinase (PI3K); protein kinase B (Akt/PKB); laminin; integrin β 1

Introduction

The association of the extracellular matrix with cell receptor proteins is central to maintaining muscle function and integrity. Defects in the protein complexes that mediate this linkage underlie the pathology in a variety of muscle diseases (Kanagawa and Toda, 2006; Lapidot et al., 2004; Rybakova et al., 2006). Several studies suggest that the structural linkage between the muscle extracellular matrix with receptors and receptor interactions with the cytoskeleton

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is crucial to prevent the progression of muscular dystrophy (Glass, 2005; Lansman and Franco-Obregon, 2006).

Cell-matrix contact is predominantly maintained through the interaction of muscle cell transmembrane receptors with laminin and other matrix proteins. The laminins, a large family of matrix proteins, are known to be multifunctional, performing key roles in development, differentiation, and migration through their ability to interact with cells via cell surface receptors (Gullberg et al., 1999). All laminin isoforms detected in skeletal muscle are recognized by $\alpha 3\beta 1$, $\alpha 6\beta 1$, and $\alpha 7\beta 1$ integrins (Fujiwara et al., 2001; Kikkawa et al., 2000). Only two laminin receptor systems are present throughout the sarcolemma of mature skeletal muscle, namely the dystrophin glycoprotein complex (DGC) and $\alpha 7\beta 1$ integrin (Mayer, 2003). Both have critical roles in the maintenance of muscle integrity (Guo et al., 2006).

The DGC provides a linkage between the cytoskeleton and the extracellular matrix. The loss-of-function mutations in DGC proteins were identified as the cause of DMD and other muscular dystrophies (Brown et al., 1999; Langenbach and Rando, 2002; Rooney et al., 2006; Watchko et al., 2002). Dystrophin's N-terminal domain binds to F-actin while its C-terminal domain binds to β -dystroglycan, which in turn interacts with the laminin-binding subunit α -dystroglycan (α DG) (Campbell, 1995). The DGC also plays a role in cell signaling (Glass, 2005; Matsumura et al., 1997; Oak and Jarrett, 2003; Zhou et al., 2005; Zhou et al., 2006). Thus, α DG may be part of a signaling pathway for the maturation and maintenance of skeletal myofibres (Brown et al., 1999; Michele and Campbell, 2003).

Integrins are heterodimeric laminin receptors that are transmembrane, link to the cytoskeleton and mediate cell-cell and cell-matrix interactions (Hynes, 2002; Rooney et al., 2006). Eighteen α and eight β chains have been identified (Gullberg et al., 1999). Among these, the $\beta 1$ integrin family forms the largest group of receptors for extracellular matrix proteins (Mayer, 2003) and all skeletal muscle integrins contain $\beta 1$. Recent studies using function-blocking antibodies have revealed the $\beta 1$ associated integrins to be important in mediating cell adhesion and differentiation (Ekstrom et al., 2003; Farias et al., 2005; Stawowy et al., 2005; Tian et al., 2002; Urbich et al., 2000). Integrins provide a similar link between the cytoskeleton and the extracellular matrix as does the DGC (Sotgia et al., 2003). Previous papers suggest that the DGC and $\alpha 7\beta 1$ integrin are independently controlled laminin receptors. Increased expression of the $\alpha 7\beta 1$ integrin has been observed in DMD patients and *mdx* mice, and raised the possibility that integrin may functionally compensate for the loss of the DGC in disease (Burkin et al., 2001; Cohn et al., 1999; Hodges et al., 1997; Vachon et al., 1997). Integrins are capable of stabilizing muscle against destruction and ameliorating the dystrophin-deficient phenotype (Mayer, 2003).

Myogenic differentiation is a highly regulated process that is controlled by multiple factors, including extracellular matrix, transmembrane receptors, and intracellular signaling molecules. Therefore, one model of the pathogenesis, which leads to cell apoptosis or necrosis in the muscle dystrophies, is through interruption of the DGC's interaction with the extracellular matrix resulting in a loss of cellular signaling (Langenbach and Rando, 2002).

The PI3K/Akt pathway is necessary to prevent apoptosis in a wide variety of cells. The PI3K/Akt pathway also has a role in the process of myotube differentiation (Ananthanarayanan et al., 2005; Glass, 2005; Lai et al., 2004; Peter and Crosbie, 2006). Activation of the PI3K/Akt signaling pathway is a key modulator of skeletal muscle hypertrophy both *in vitro* and *in vivo* (Takahashi et al., 2002). $G\beta\gamma$ can activate PI3K following binding of GTP or cholera toxin (CT) (Brock et al., 2003; Gilman, 1987; Schnitzler et al., 2007), and thereby initiate $G\beta\gamma$ mediated signal transduction pathways. Activation of PI3K and formation of its lipid products *in vivo* lead to activation of Akt and downstream inhibition of glycogen synthase kinase-3

(GSK-3), which are involved in cell survival and protein synthesis pathways (Baar and Esser, 1999; Pap and Cooper, 1998).

An understanding of the relevant signal transduction pathways and of the interactions between these pathways in the skeletal muscle cell will facilitate efforts to elucidate the pathogenesis of muscular dystrophies. To understand the role of the PI3K-Akt signaling in muscular dystrophies, we perform a detailed analysis of the protein interactions between the DGC and PI3K/Akt signaling in skeletal muscle, and then we investigated the role of G β γ -dimers, laminin and its receptors in the activation of PI3K/Akt. We also investigated whether perturbation of these interactions could lead to the disruption of PI3K/Akt signaling in muscle cells. The results demonstrate the existence of a specific link between the laminin-DGC-G β γ -PI3K-Akt signaling in skeletal muscle. G β γ binding activates PI3K/Akt signaling in a laminin-dependent manner, and phosphorylation of Akt and GSK result from activation of PI3K. This reveals further details of how the PI3K/Akt pathway becomes activated upon binding of the DGC to the extracellular matrix. This laminin-DGC-G β γ -PI3K-Akt signaling is likely to be important to the pathogenesis of muscular dystrophies. Up-regulating integrin β 1 expression and its signaling may partially compensate for the loss of dystrophin in *mdx* mice.

MATERIALS AND METHODS

Materials

Rabbit antibodies against G β , PI3Kp110, Akt1, Akt 1/2, actin (C-2), Na⁺,K⁺-ATPase (H-300) and integrin β 1 and mouse monoclonal antibodies against PI3Kp85 and pAkt were from Santa Cruz Biotechnology Inc. (California, USA). PI3K inhibitors LY294002, wortmannin and rabbit polyclonal antibodies against phospho-GSK3 β were purchased from Cell Signaling Technology, Inc. (Beverly, MA, USA). GTP- γ S, GDP and cholera toxin were from Sigma Chemical, Inc. (St. Louis, Mo, USA). Mouse laminin-1 was from BD Biosciences, Inc. (Bedford, MA, USA). Mouse monoclonal α DG antibodies VIA4-1 and I1H6 were the generous gifts from Dr. Kevin P. Campbell (University of Iowa, USA). Affinity purified rabbit polyclonal antibody against β -dystroglycan was a generous gift from Dr. Tamara C. Petrucci (Laboratorio di Biologia Cellulare, Istituto Superiore di Sanita, Via le Regina Elena, Roma, Italia). Goat anti-mouse IgG (H+L)-horseradish peroxidase conjugate and goat anti-rabbit IgG (H+L)-horseradish peroxidase conjugate were from Bio-Rad Laboratories, Inc. (USA). Phosphatidylinositol (cat. # P0639) and phosphatidylinositol-3-phosphate (P0474) and histone (H4524) were from Sigma. γ -³²P-ATP was from MP Biomedical. The mouse C2C12 cell line was obtained from the American Type Culture Collection (Rockville, MD). All other chemicals were of the highest purity available commercially.

Preparation of Microsomes from Rabbit Skeletal Muscle

Frozen rabbit skeletal muscle (3 g back muscle) was homogenized in 21 ml pyrophosphate buffer (20 mM sodium pyrophosphate, 20 mM phosphate, 1 mM MgCl₂, 0.303 M sucrose, and 0.5 mM EDTA, pH 7.0) in the presence of a cocktail of protease inhibitors. The homogenate was centrifuged at 13000 g for 15 min at 4°C. The supernatant was then centrifuge for 30 min at 32500 g to pellet muscle microsomes. The pellets were suspended in 0.6 ml buffer A (50 mM Tris, pH 7.5, 100 mM NaCl).

Laminin Depletion

Laminins were depleted from skeletal muscle membrane using heparin-Sepharose as previously described (Oak and Jarrett, 2003). Rabbit muscle microsomes were incubated with either Sepharose 4B as a negative control (retaining endogenous laminins) or heparin-Sepharose (to deplete laminins) for 1 h at 4°C, with gentle mixing. After incubation, the resin

was removed by slow speed centrifugation (2000 rpm) and the supernatant microsomes were used. Microsome were adjusted to equal protein concentration.

Bovine brain Gβγ was purified as described (Katada et al., 1986; Zhou et al., 2005).

PI3K assays

Immunoprecipitates with either PI3K-p110 or VIA4 antibodies were prepared and mixed with purified Gβγ and added to a PI3K assay previously described (Dogra et al., 2006). The assay uses phosphatidylinositol as substrate and γ -³²P-ATP to follow the production of phosphatidylinositol-3-³²PO₄, separated by silica thin layer chromatography and detected by autoradiography. Phosphatidylinositol and phosphatidylinositol-3-PO₄ were also run on the same plate and stained (Dittmer and Lester, 1964) to determine the Rf for substrate and product, respectively.

Akt assay

Immunoprecipitates with either Akt1/2 or VIA4 antibodies were prepared and mixed with purified Gβγ and added to an Akt assay previously described (Dogra et al., 2006) using histone H2B as substrate.

Immunoprecipitation

Rabbit muscle microsomes were isolated and incubated for 30 min in buffer A containing 1mM CaCl₂, 1mM MgCl₂, 1mM GTP- γ S, 1mM ATP and with or without 3 μ g laminin-1. After pre-clearing with protein G-Sepharose, 5 μ g of specific antibodies was added and incubated for 1 h at 4°C with gentle mixing. The membranes were solubilized in 1% digitonin, and incubation was continued for another 1 h. The immunoprecipitates were collected by adding protein G-Sepharose, and incubating for 2 h at 4°C. After washing with buffer K (20 mM Hepes, pH 7.5, 10 mM MgCl₂, 100 mM KCl), bound protein was eluted with SDS-PAGE sample buffer.

Western Blotting

Samples were separated by 12% SDS-PAGE, and transferred to nitrocellulose membranes at 4°C. Membranes were blocked with 5% nonfat milk and incubated with the indicated primary antibodies. Primary antibodies VIA4, PI3Kp85, pAkt, Akt, p-GSK-3 β , integrin β 1 and β -dystroglycan were used at a 1:1000 dilution. Immunoreactive bands were then detected with horseradish peroxidase-conjugated secondary antibodies. Anti-rabbit IgG was diluted 1:10000, and anti-mouse IgG were used at a 1:5000 dilution. Membranes were then stripped by incubation at 50°C in 2% SDS, 100 mM 2-mercaptoethanol, 62.5 mM Tris, pH 6.8. The blots were then blocked again with milk and probed with additional primary antibodies, typically to provide a loading control. Blots were developed using enhanced chemiluminescence (ECL) as previously described (Oak and Jarrett, 2003) and exposure to X-ray film.

Cell Culture

Mouse C2C12 myoblasts were maintained in Dulbecco's modified Eagle's medium (DMEM) supplemented with 10% fetal bovine serum (FBS) at 37°C in a humidified atmosphere of 5% CO₂ in air. Differentiation to myotubes was initiated at 80% confluence by replacing the growth medium with DMEM containing 1% FBS and maintaining the cells in this medium for 5 days. On the fifth day, myotubes were washed with PBS and harvest by adding 1 ml of trypsin-EDTA solution (0.05% trypsin, 0.02% EDTA in Hank's Balanced Salt Solution, HBSS) for 5 min at 37°C. The cells were then immediately suspended in DMEM containing 10% FBS and removed from the plate. Myotubes (10⁶/ml, 1 ml) were cultured in six well plates for 24 h in the absence or presence of increasing concentrations of LY294002 or wortmannin. At the termination of the inhibition experiment, cells on the plate were washed with PBS and used for the MTS assay

of cell viability using the CellTiter 96[®] AQueous reagent and protocols (Promega, Madison WI) or were then lysed in 0.3 ml/well of modified RIPA lysis buffer (1% Igepal CA-630, 0.5% sodium deoxycholate, 0.1% SDS in PBS) supplemented with protease inhibitors (1 µg/ml pepstatin A, 5 µg/ml aprotinin, 1 µg/ml leupeptin, 0.75 mM benzamidine, and 0.1 mM phenylmethylsulfonyl fluoride). Following incubation for 30 min at 4°C, cell lysates were clarified by centrifugation at 15000 g for 15 min, the supernatant mixed with SDS-PAGE sample buffer, and Western blotted.

In other experiments, myotubes were cultured in the presence of 10 µg/ml of VIA4 or IIIH6 antibodies and microsomes prepared from the myotubes (Fig. 2C).

Preparation of Laminin- Sepharose

CNBr-preactivated Sepharose (Pharmacia) was suspended in ice-cold 1 mM HCl for 15 min and washed with 1 mM HCl on a sintered glass filter. Laminin-1 (1 mg) in coupling buffer (0.1 M NaHCO₃, 0.5 M NaCl, pH 8.3) was mixed with 0.5 ml of the activated Sepharose in a screwcap plastic tube on a wheel rotator mixer overnight at 4°C. The laminin-Sepharose was washed with blocking buffer (0.1 M Tris-HCl, 0.5 M NaCl, pH 8.0), and mixing continued in this buffer overnight. The resin was stored in blocking buffer containing 10 mM NaN₃ as a 1:1 slurry.

Pull-Down Assay

Laminin-Sepharose (100 µL of a 50% slurry) was washed with buffer K containing 1 mM CaCl₂. Microsomes in buffer K containing 1 mM CaCl₂, 1 mM GTP-γS and 1 mM ATP were added and incubated for 1 h at 4°C. The microsomes were solubilized by addition of digitonin to 1% and incubation continued for 1 h at 4°C. The samples were centrifuged for 1 min at 2000 rpm in a microcentrifuge. The resins were washed three times with buffer K. Bound protein was eluted with 60 µL of twice concentrated SDS-PAGE sample buffer. Samples were heated for 5 min at 95°C and centrifuge for 5 min to remove the resins. Supernatants were applied to 12% SDS-PAGE and electroblotted to nitrocellulose. The blots were blocked with 5% nonfat milk in TTBS (20 mM Tris, 0.1 M NaCl, 0.2% Tween-20, pH 7.5). Blots were incubated with 1:1000 diluted antibodies against integrin β1 or β-dystroglycan. Goat anti-rabbit IgG (H+L)-horseradish peroxidase conjugate (1: 10000) was used as a secondary antibody. The blots were developed using ECL.

Animal Models

Mdx and wild-type (C57BL/6) breeder pairs were purchased from Jackson Laboratories (Bar Harbor, ME). Mice were housed in the light- and temperature-controlled quarters where they received food and water *ad libitum*. Mice were anesthetized with isoflurane for surgery and tissue removal follow by exsanguination. To examine *mdx* disease progression, four mice were analyzed at 4 weeks, and six adult mice were analyzed at 6 month of age. Skeletal muscles (gastrocnemius) were harvested and snap frozen in liquid nitrogen. All tissues were stored at -80°C until analysis. All procedures were carried in accordance with guidelines set by the UTSA Institutional Animal Care and Use Committee.

Cholera Toxin Experiment

C2C12 myoblasts (10⁶/ml, 1 ml) in DMEM medium supplemented with 10% fetal bovine serum were placed in a 24 well plate and maintained at 37°C and 5% CO₂ for 30 min in a humidified incubator. Cholera toxin was added to 100 ng/ml for 0, 5, 10, and 30 min. The cells were then washed with PBS, and dissolved in RIPA buffer as previously described. The clarified cell lysates were separated on 12% SDS-PAGE and transferred to nitrocellulose

membranes. The membranes were probed with antibodies against phosphorylated Akt followed by ECL. The membranes were then stripped as previously described and probed for total Akt.

Loading Controls

Several specific loading controls are specified in the figure legends. All blots were stained for protein with Ponceau Red (0.2% Ponceau Red, 3% trichloroacetic acid, 3% 5-sulfosalicylic acid), a reversible stain, prior to blocking with milk to observe whether loading was equal. Also blots were usually stripped after the first western blot, western blotting was repeated with a different primary antibody to detect an unrelated protein, typically β -actin, total Akt or β -dystroglycan, confirming equal loading. For each experiment, 2–3 different exposures of each film were taken. In most cases, a longer exposure time is shown in the figures for clarity but shorter exposure time films were used for the densitometry analysis reported in the text, since these are less prone to film saturation errors. In addition, each experiment was repeated at least 3 times.

Statistical

Unless otherwise stated, densitometry was performed on two independent experiments where the ECL or autoradiographic films were underexposed, to avoid problems with film saturation. Data are reported as the mean \pm standard deviation. Statistical significance was assessed using Student's t-test (1 tail with unequal sample variance).

RESULTS

Interactions between the DGC-G $\beta\gamma$ and the PI3K /Akt pathway

For most of the experiments here, we use antibodies from different species (e.g., mouse and rabbit) for immunoprecipitation and western blotting, which allows us to perform western blots without the interference of observing immunoglobulin heavy chain. This allows us to observe Akt (60 kDa.) clearly.

Previously, we had shown that heterotrimeric G $\alpha\beta\gamma$ was bound by DGC syntrophin in a laminin-independent manner (Zhou et al., 2005). To examine whether the DGC, G $\beta\gamma$, PI3K (the p85 or p110 subunits) and Akt form a complex in skeletal muscle, four different immunoprecipitation experiments were performed, and then interactions were analyzed by western blot. The results show that regardless of whether antibodies against Akt (Fig. 1A), G β (Fig. 1B), PI3K (Fig. 1C), or DGC α -dystroglycan (Fig. 1D) are used for immunoprecipitation, the other components are detected by western blots, showing that all are associated with one another. These results confirm that the DGC is associated with several signaling molecules, including G $\beta\gamma$, PI3K and Akt, which are co-localized at the skeletal muscle membrane. In Fig. 1A–D, an equivalent amount of the crude microsomes (without immunoprecipitation) is also shown. Essentially all of the microsomal pAkt immunoprecipitate and much of the α -dystroglycan, PI3K, and total Akt immunoprecipitate in this complex.

As negative controls, similar experiments were also performed with pre-immune IgG or with the Na⁺, K⁺-ATPase antibody since Na⁺, K⁺-ATPase is also a membrane protein but with no known association with these cell signaling components. The results in Fig. 1E show that while the detected proteins are present in the crude microsomes, they are not immunoprecipitated.

The role of G $\beta\gamma$ and laminin in the activation of PI3K

Heterotrimeric G-proteins consist of a G α -subunit and G $\beta\gamma$ -dimer. The G α -subunit binds guanine nucleotides (GDP or GTP) and possesses an intrinsic low turnover number GTPase activity (Schwindinger and Robishaw, 2001). G $\beta\gamma$ always remains tightly associated as a

heterodimer. The $G\beta\gamma$ -dimer does not dissociate from $G\alpha$ under physiologic conditions except when the $G\alpha$ -subunits are in the GTP-bound state. Both $G\alpha$ and $G\beta\gamma$ initiate intracellular signaling through a variety of effector molecules (Maier et al., 2000), but in many cases it is the $G\beta\gamma$ rather than the $G\alpha$ that transmits signals for proliferation and survival (Schwindinger and Robishaw, 2001). Here we will focus on the role of $G\beta\gamma$ in the activation of PI3K, an important step in G-protein coupled survival signaling. We investigated whether this Akt activation results from $G\beta\gamma$ using two experiments. In Fig. 2A, rabbit skeletal muscle microsomes were immunoprecipitated with an antibody against α DG (VIA4) in the presence of either 1 mM GTP- γ S or 1 mM GDP. GDP should force the $G\alpha\beta\gamma$ state and inhibit PI3K while the non-hydrolysable GTP- γ S analogue, should have the opposite effect (Klein et al., 2000; Xie et al., 2000). PI3K activity was then monitored by down-stream Akt activation using the phosphorylated pAkt (Ser 473) antibody. The blot was then stripped and re-probed with an antibody to actin (known to be associated with the DGC) or total Akt 1/2 (data not shown), as loading controls. As expected interactions between α DG and pAkt were higher in the presence of GTP- γ S compared to GDP, while actin (Fig. 2A) was not significantly altered. Based upon densitometry, GTP- γ S gave 4.3-fold higher pAkt activation while loading on the two blots assessed by β -actin differed by less than 4% (Fig. 2A). For other panels in the figure, C2C12 myotubes were used to measure the effect of inhibitors and a blocking antibody. Wortmannin (200 nM), an inhibitor of PI3K, prevents the pAkt activation, showing that this activation requires PI3K (Fig. 2B) while the total amount of Akt is unaltered Wortmannin (200 nM) resulted in 6.2-fold lower pAkt (6.2 ± 0.7 , $n=2$), while total Akt differed by less than 3% (Fig. 2B).

To further probe whether laminin-binding to the DGC is required, an additional α DG antibody, I1H6 was included in Fig. 2C. The I1H6 antibody binds α DG and blocks its interaction with laminin while the VIA4 antibody, used in other experiments, does not (Ervasti and Campbell, 1993). When immunoprecipitation is performed with either antibody, β DG is found showing that both immunoprecipitate the DGC, but with the blocking antibody, there is no longer an association of the DGC with PI3K detected. By densitometry, there is 243-fold more PI3K with the non-blocking VIA4 antibody than there is with the blocking I1H6 antibody, while both antibodies precipitate nearly equal amounts of β DG, differing by less than 20%. These results strongly suggests that the laminin-DGC interaction is required for the DGC- $G\beta\gamma$ -PI3K-Akt signaling.

The heterotrimeric G-protein associated with the DGC is G_s (Zhou et al., 2005) and the GTPase activity of G_s is irreversibly inhibited by cholera toxin; thus, both GTP- γ S and cholera toxin would increase the dissociation of $G\alpha$ from $G\beta\gamma$. This results in more free $G\beta\gamma$ for activation of PI3K and Akt. As shown in Fig. 2D, incubation for 5–30 min with 100 ng/ml cholera toxin caused an increase in the level of phosphorylated pAkt compared to control (upper panel) while total Akt (lower panel) provides a loading control. By densitometry, at 10 min, pAkt increased 4.9-fold (± 0.4 , $n=2$) relative to no cholera toxin. These results confirm the results with nucleotides that downstream Akt is activated via $G\beta\gamma$. Interestingly, wortmannin and cholera toxin affect the activation of Akt to pAkt without altering its total amount or association with the DGC or in the C2C12 myotubes, respectively.

Laminin-binding increases PI3K/Akt signaling

Laminins are multifunctional, performing key roles in development, differentiation, and migration through their ability to interact with cells via cell surface receptors, including α -dystroglycan. In adult skeletal muscle, laminin-2 ($\alpha_2\beta_1\gamma_1$) is the major laminin present endogenously while laminin-1 ($\alpha_1\beta_1\gamma_1$) is known to have many of the same effects. Whether laminins affect G-protein binding and activates PI3K/Akt signaling via the DGC is not well understood but is strongly suggested by the results in Fig. 2C. Previously (Oak and Jarrett,

2003), we showed that laminins can be depleted from skeletal muscle microsomes and here we used laminin-depletion to examine the phosphorylation state of pAkt and the amount of total Akt bound (Fig 3A). When endogenous (endo) laminin is present, Akt is activated (pAkt). When laminin is depleted (-), less active pAkt is found, 50% (± 11 , $n=2$) less by densitometry, and activation is restored to 92% of the endogenous laminin amount by the addition of exogenous (exo) laminin-1. Thus, either the endogenous laminin-2 or exogenous laminin-1 have similar effects. The same membrane was stripped and detected for total Akt which is unaffected by these treatment (Fig. 3A), thus Akt is associated in all cases but only activated to pAkt when laminin is present. GDP, which prevents G $\beta\gamma$ release, prevents the laminin-dependent activation of Akt (Fig. 3B), showing this proceeds through a G-protein. Indeed, pAkt is so low, it is difficult to see in the figure while total Akt is clearly observed.

To observe this GTP- γ S/GDP effect more clearly, in an independent experiment samples were applied to the same gel, blotted, and pAkt detected by Western Blot (Fig. 3C). The results show pAkt is present in crude microsomes in an amount similar to that seen after GTP- γ S treatment and PI3K-immunoprecipitation. Thus, most pAkt present in microsomes must be associated with PI3K in a complex. Furthermore, untreated microsomes behave as though they are in a GTP-rich environment, not surprising given the prevalence of GTP over GDP in muscle and essentially all tissues. GDP-treatment results in such low amounts of pAkt immunoprecipitating with PI3K that it is not observed at this exposure (although longer exposures do reveal a small amount). Again, laminin-depletion affects pAkt activation.

We also investigated the effects of laminin on the interactions of DGC-G $\beta\gamma$ -PI3K-pAkt in Fig. 3D. The data show that regardless of whether immunoprecipitation is with DGC α -dystroglycan or G β , PI3K and pAkt co-precipitate and that these components are greater in the presence of either endogenous or exogenous laminin than in laminin-depleted microsomes. Since total Akt remains associated regardless of laminin, laminin must be required for its activation to pAkt but not its association with the complex. These findings strongly suggest that laminin binding can initiate PI3K/Akt cell signaling. DGC-G $\beta\gamma$ is binding PI3K and pAkt is activated in a laminin-dependent manner. Densitometry results confirm what is seen by visual inspection. In the upper panel of Fig. 3D, PI3Kp85 decreases by $71\pm 31\%$ upon depletion of laminin, while in the middle panel, pAkt decreases $68\pm 33\%$, and in the lower panel, PI3Kp110 is decreased by $37\pm 9\%$ upon depletion of laminin.

Purified bovine brain G $\beta\gamma$ activates PI3K and pAkt and these are associated with the DGC

Fig. 4A shows that the addition of purified G $\beta\gamma$ increases the enzymatic activity of PI3K. Densitometry of independent experiments confirms that the activation is increased 24 ± 1 -fold when $6\ \mu\text{g/ml}$ G $\beta\gamma$ is used and the activation is dose-dependent (Fig. 4B). The PI3K associated with the DGC α -dystroglycan was also increased by G $\beta\gamma$ as shown in Fig. 4C. In this latter figure, a longer exposure is shown and just the upper region of the chromatogram is shown to emphasize a second, slower migrating phospholipid also observed. Since PI3K can phosphorylate microsomal phosphatidylinositides at the 3-position of inositol, it is likely this additional spot is microsomal lipid which differs from the added substrate by either having different fatty acyl chains or additional phosphorylations on the inositol at other positions (e.g., 4,5). The latter possibility would be consistent with slower migration on the silica thin layer chromatogram.

Figure 5A shows that purified G $\beta\gamma$ also increases the DGC associated activated pAkt. Syntrophin provides a loading control. Based on densitometry of two independent experiments, for Fig. 5A, G $\beta\gamma$ increased pAkt 4.3 ± 2.8 -fold while syntrophin was unchanged (1.05 ± 0.03 -fold). Presumably, this pAkt activation is a result of the increased PI3K activity shown in Fig. 4 since it is inhibited by the PI3K inhibitor wortmannin (Fig. 2B, and see below). That this pAkt is enzymatically activated is shown in Fig. 5B, where G $\beta\gamma$ activates the phosphorylation of

histone H2B, a known substrate for pAkt (Dogra et al., 2006). Based on densitometry of two independent experiments, for Fig. 5B, G β increased pAkt activity 3.3 ± 1.9 -fold and 4.8 ± 3.2 -fold at 2 and 6 μ g/ml. respectively.

Thus, we conclude that the form of PI3K present in muscle and which associates with the DGC is activated by G β . This is also consistent with the effect of cholera toxin, guanine nucleotides and α -dystroglycan blocking antibodies already shown (Fig. 2 and 3) and that this PI3K is resulting in pAkt activation.

Effect of Wortmannin on PI3K activation

Activation of PI3K and formation of its lipid products *in vivo* lead to activation of pAkt, the downstream serine-threonine protein kinase involved in cell survival and protein synthesis pathways (Li et al., 2005). Whether this activation of Akt or downstream GSK is occurring through PI3K was next further investigated. Two pharmacological inhibitor of PI3K, LY294002 and Wortmannin, were added to cultured C2C12 myotubes. Cultured myotubes were incubated for 24 h with LY294002 at concentration from 25 μ M to 150 μ M (data not shown) or Wortmannin at concentration 100 nM or 200 nM (Fig. 6A–D) and the phosphorylation state of Akt (Fig. 6A and B) and GSK-3 β (Fig. 6C and D) were examined in treated and untreated myotube cultures. pAkt and pGSK-3 β phosphorylation were reduced by either PI3K inhibitor, and the inhibitory effects of Wortmannin were more pronounced than the effects of LY294002 over these concentrations and is shown in Fig. 6. Densitometry results are shown in Fig. 6B and D. The amounts of total Akt and actin were unchanged, providing loading controls. These results strongly suggest that the phosphorylation of pAkt or pGSK is mediated *via* PI3K.

One concern is the effect of such inhibitors on cell viability. Shown in Fig. 6E is a measure of cell viability for the two inhibitors, using the MTS assay. While Wortmannin at 200 nM has only minimal effect on viability and cannot account for the changes seen in Fig. 6A–D, LY294002 decreases the number of viable cells by over 50%. Thus, the Wortmannin results on cell signaling are not caused by an effect on cell viability while the results with LY294002 are more problematic and are not shown.

PI3K/Akt signaling is upregulated in skeletal muscle of *mdx* mice

The *mdx* mouse has a dystrophin mutation which makes it severely deficient in dystrophin and in most of the components of the DGC (Ohlendieck and Campbell, 1991). Others have previously reported that total Akt is elevated in *mdx* mouse at 12 weeks of age (Dogra et al., 2006). In order to investigate how the loss of the DGC impacts PI3K/Akt signaling in the *mdx* mouse model of Duchenne muscular dystrophy, we examined pAkt phosphorylation and total Akt levels in the gastrocnemius muscle from *mdx* mice and their normal control (C57bl/6) in Fig. 7. *Mdx* mice exhibit a significant increase ($p < 0.05$, $n = 2$) in pAkt compared to age-matched normal mice, but total Akt (Akt 1 and 2) is not affected (Fig. 7A). By densitometry, *mdx* mouse at this age have 2.1 ± 0.9 -fold greater pAkt (two independent experiments) while total Akt is insignificantly decreased by 7%. Similar results were also obtained in 6 month and older adult mice for the gastrocnemius (data not shown). pGSK was not changed between normal and *mdx* mice (Fig. 7B). One possible explanation for why pAkt could become more active, while its downstream substrate GSK could remain constant would be if GSK is completely phosphorylated even at the lower activity of pAkt found in normal mice. Thus, *mdx* displays elevated pAkt but similar amounts of total Akt and pGSK compared to normal tissue.

Enhanced expression of integrin $\beta 1$ and $G\beta$ in *mdx* skeletal muscle induces activation PI3K/Akt

Muscle fibers attach to basal lamina laminin using either the dystrophin glycoprotein complex or integrins (Burkin et al., 2001). Thus, both the integrin- and dystrophin-mediated transmembrane linkage contribute to the functional integrity of skeletal muscle (Guo et al., 2006), and defects in either linkage system results in muscular dystrophy (Burkin et al., 2001). Our data and those of others (Dogra et al., 2006) show the PI3K/Akt pathway was upregulated in *mdx* mice compared to normal tissue. This has led us to hypothesize that regulation of pAkt is defective in *mdx* mice and may proceed through an integrin-mediated process rather than a DGC-mediated one. To test this hypothesis we analyzed the expression of integrins and DGC β -dystroglycan in the gastrocnemius muscle. Since all muscle integrins contain the integrin $\beta 1$ subunit, this was analysed to reflect total integrin expression. The results in Fig. 8 show that integrin $\beta 1$ is expressed at a much higher level in *mdx* muscle relative to control (Fig. 8A) while β -dystroglycan was greatly reduced (Fig. 8B) as expected (Ohlendieck and Campbell, 1991). Based on densitometry, integrin increased 4.0 ± 0.2 -fold ($n=2$) while β -dystroglycan decreased 4.8-fold in *mdx* mouse relative to control ($n=2$). In Fig. 8C and D, we demonstrate a higher interaction between integrin $\beta 1$ with both laminin and PI3K in *mdx* mouse microsomes relative to control in vitro. By densitometry, 7.2 ± 5.1 -fold ($n=2$) as much integrin was co-immunoprecipitated with PI3Kp85 in *mdx* mouse (Fig. 8C) while at least 3-fold greater amounts of integrin were pulled down by laminin-Sepharose in *mdx* mouse gastrocnemius ($n=2$). Blots were stripped and probed for actin to provide a control for equal loading (data not shown). In Fig. 8E, the results show that $G\beta$ was increased in *mdx* muscle microsome compared with control (by densitometry, 2.2 ± 0.1 -fold, $n=2$) and interactions of PI3K- $G\beta$ (11.1 ± 1.3 -fold, $n=2$) and integrin $\beta 1$ - $G\beta$ (7.9 ± 1.4 -fold, $n=2$) were also enhanced in *mdx* mice. These results suggest that PI3K signaling was increased due to the increased expression of integrins and $G\beta$ as well as their increased interaction with laminin and PI3K in *mdx* mice (Fig. 8C and D), which may partially compensate for the absence of the dystrophin and diminished amounts of other DGC components (Farias et al., 2005; Wang et al., 2007). These results further suggest that disruption of the DGC in *mdx* mouse is causing dis-regulation of the laminin-DGC- $G\beta\gamma$ -PI3K-Akt signaling, which is likely being replaced by integrin- $G\beta\gamma$ -PI3K-Akt signaling.

DISCUSSION

The results show that the DGC, $G\beta\gamma$, PI3K, and Akt exist as a complex in skeletal muscle microsomes (Fig. 1). Our previous work had shown that the Gs heterotrimeric G-protein bound to the DGC syntrophin via syntrophin's PDZ domain and that the activation state of Gs was laminin-dependent. The $G\alpha$ apparently interacts with the L-type Ca^{2+} -channels to lessen intracellular Ca^{2+} when laminin is bound (Zhou et al., 2005). Possible $G\beta\gamma$ effects were not then explored and are the subject of the current investigation. Laminin-binding to the DGC also causes activation of pAkt and downstream signaling which prevents apoptosis (Langenbach and Rando, 2002). Upstream of pAkt is PI3K. Here, we have presented convincing evidence that this DGC-associated $G\beta\gamma$ associates with PI3K, activates it, and causes the pAkt activation.

PI3Ks in mammals are the products of at least eight different genes divided into three different classes (Hirsch et al., 2007). The combination of antibodies used here would be consistent with the PI3K type 1A while most of the literature on $G\beta\gamma$ activation of PI3K involves members of the type 1B PI3Ks. However, there are clear reports of PI3K type 1A being activated by $G\beta\gamma$ (Hirsch et al., 2007; Murga et al., 2000). Type 1A PI3Ks are typically recruited to the membrane by interactions between PI3Kp85 SH2 domains with phosphotyrosine sequences on a target protein and are also activated by Ras (Hirsch et al., 2007). Since syntrophin is phosphorylated by Src family tyrosine kinases (Zhou et al., 2007; Zhou et al., 2006) as has also been reported

for β -dystroglycan (Sotgia et al., 2003), recruitment to phosphotyrosine sequences may indeed be involved here, although we did no experiments to directly address this. However, this would not explain the direct effects of purified $G\beta\gamma$ on PI3K (Fig. 4) and Akt (Fig. 5). However, the PI3K class 1A p110 β subunit is also activated by $G\beta\gamma$ (Kurosu et al., 1997). It is most likely that it is this PI3K which accounts for our results.

Other G-proteins, including the small G-proteins, also activate PI3K. Activation by Ras can probably be excluded. Laminin-Sepharose pulls down a complex containing DGC proteins and Rac1 (Oak and Jarrett, 2003; Zhou et al., 2007; Zhou et al., 2005) but not Ras or other p21 G-proteins (Oak and Jarrett, 2003). Laminin-binding is required for recruiting Gs to the DGC (Zhou et al., 2005) and here we show that a specific antibody (IIH6), which blocks laminin-binding by DGC α -dystroglycan, blocks the recruitment of PI3K to the DGC (Fig. 2C). This same antibody has also been shown to block $G\beta\gamma$ -binding (Zhou et al., 2005) and Akt activation and to induce apoptosis (Langenbach and Rando, 2002) in muscle cells. Thus, the data are most consistent with either Rac1 or $G\beta\gamma$ being the G-protein responsible. However, cholera toxin should specifically affect only Gs in a way that would increase the availability of $G\beta\gamma$ as was shown in Fig. 2E by its effect on the activation of pAkt, and the direct effect of purified $G\beta\gamma$ confirms $G\beta\gamma$. Thus, the results are consistent with a model in which laminin-binding to the DGC increases the availability of $G\beta\gamma$ and this is in turn activating Type 1A PI3K, probably containing the p110 β catalytic subunit.

Laminins interact with cell surface receptors, leading to intracellular signaling and cytoskeletal reorganization, and thus have important regulatory roles. Laminin binding to α -dystroglycan or integrins can initiate cell signaling (Hayashi et al., 2001; Miyagoe et al., 1997). However, the role of laminin in the activation of PI3K is unclear and has been a focus of this investigation. The result demonstrates that phosphorylated pAkt was reduced as a result of laminin depletion using heparin-Sepharose, but adding exogenous laminin to the depleted microsomes reconstitutes pAkt activation (Fig 3A). To elucidate further the mechanism of the laminin induce signaling we investigated the effects of laminin on interaction of DGC- $G\beta\gamma$ -PI3K-Akt (Fig 3B). The data shows laminin increases binding activity, and laminin-deficiency decreases the interactions of DGC- $G\beta\gamma$ -PI3K-pAkt. Either endogenous laminin (predominantly laminin-2) or exogenous laminin-1 can enhance signaling. These findings suggest that DGC- $G\beta\gamma$ is binding PI3K and activating pAkt in a laminin-dependent manner. Taken together with previous results, which show that the same IIH6 antibody that blocks the laminin-dystroglycan interaction, inhibits Akt signaling and initiate apoptosis (Langenbach and Rando, 2002), leaves little doubt that this DGC- $G\beta\gamma$ -PI3K-Akt signaling is an important pathway for muscle cell survival.

To determine whether this phosphorylation of Akt or downstream GSK is occurring through PI3K, we have investigated the effects of two pharmacological inhibitor of PI3K on Akt and GSK phosphorylation in cultured C2C12 myotubes. The results show that 150–200 nM wortmannin can inhibit activation of Akt and GSK with little decrease in cell viability (Fig 4), and confirm the PI3K/Akt pathway play an important role mediating cell–survival signaling. LY294002 was also an effective inhibitor but its effects on myotube viability makes the meaning of these results much less clear.

How this may function in intact muscle, where laminin is stably bound, was not investigated. A recent paper (Zhou et al., 2007), however, suggests how this may occur. Laminin-binding to the DGC also results in signaling through a DGC/Rac1/JNK pathway. In this case, stretching or contracting intact muscle also resulted in signaling and it was proposed that it is stressing the laminin-DGC linkage during muscle activity is what initiates signaling *in vivo*. The same may apply here.

In order to investigate how the loss of the DGC impacts PI3K/Akt signaling during *mdx* dystrophy we investigated Akt activation in normal and dystrophic muscle. The results show that pAkt levels were significantly elevated in *mdx* skeletal muscle relative to control in agreement with a previous report (Dogra et al., 2006). These data showing the activation of PI3K/Akt signaling in dystrophin-deficient muscle (Fig. 7 and 8) suggesting that the response of *mdx* muscle is initiated early in the pathogenic process (Peter and Crosbie, 2006). Akt activation was detected at 4 weeks, at a very early stage of disease pathogenesis as well as at later stages (data not shown). DMD patients exhibit a similar pattern of Akt activation (Dogra et al., 2006; Peter and Crosbie, 2006). Recent investigation have demonstrated the importance of the PI3K/Akt pathway in regulation of muscle apoptosis (Rommel et al., 2001). In our data 4-week-old *mdx* mice display elevated Akt signaling suggesting a compensatory cell survival response to the disease.

Recently, studies demonstrate that there is an increase in the amount of integrin in DMD patients and *mdx* mice, which could ameliorate the development of muscle diseases (Burkin et al., 2001; Li et al., 2003), but the mechanism is not currently understood. It remains possible that compensatory mechanisms may be activated via alternative receptor systems such as integrins and may compensate for defective DGC/PI3K/Akt signaling mechanisms (Burkin et al., 2005). Our results demonstrate not only increased integrin $\beta 1$ expression, but also significantly increased the interaction between integrin $\beta 1$, laminin, $G\beta\gamma$ and PI3K occurs in *mdx* mice (Fig. 8). These results point to a mechanism by which enhanced levels of integrin $\beta 1$ activate the $G\beta\gamma$ /PI3K/Akt pathway in muscle (Farias et al., 2005). Integrin $\beta 1$ may compensate for the defects of the DGC in *mdx* mice (Burkin et al., 2005; Burkin et al., 2001; Deconinck et al., 1997) by reinforcing the signal transduction pathways which normally link the DGC to cell survival signaling.

Most G-protein coupled receptors which interact with the heterotrimeric G-proteins are of the seven transmembrane spanning receptor motif. Clearly, while the DGC is a G-protein coupled receptor, it is of a different type. In this case, the DGC is a multiprotein complex, binding laminin via the dystroglycan proteins, spanning the sarcolemma with the β -dystroglycan subunit and the sarcoglycans, and interacting with the cytoskeleton via dystrophin and G-proteins via syntrophin. The DGC is a G-protein coupled laminin-receptor. When laminin binds, $G\beta\gamma$ apparently activates PI3K, which in turn causes pAkt activation. We have previously shown that the particular region of laminin, the laminin globular (LG) domains 4–5, which binds to DGC α -dystroglycan (Talts et al., 1999), causes proliferation in myoblasts (Zhou et al., 2007) and others have shown that laminin is necessary for normal muscle differentiation and growth. Thus, if $G\beta\gamma$ becomes available, either because of laminin-binding or by other means (e.g., muscle activity), a hypertrophic response mediated by the Akt pathway would be expected. This has been observed with pharmacological agents.

$G\beta\gamma$ can also be made available at the sarcolemma by other means. Of the more typical type of G-protein receptors are the β -adrenergic receptors, of which $\beta 2$ -adrenoceptor is the most abundant muscle sub-type. This receptor also couples through heterotrimeric G-proteins (Lynch et al., 2007). $\beta 2$ - Adrenoceptor agonists, such as clenbuterol, which would also increase available sarcolemma $G\beta\gamma$, cause muscle hypertrophy and increase muscle mass. Furthermore, clenbuterol also causes increased pAkt activation and downstream signaling through mTOR and S6 kinase. The authors of these studies hypothesized that it could be the $G\beta\gamma$ released by the $\beta 2$ -adrenoceptor in response to clenbuterol that was activating PI3K/Akt to cause muscle hypertrophy (Kline et al., 2007). Here, we provide evidence this occurs for the laminin-DGC- $G\beta\gamma$ -PI3K-pAkt signaling.

There is no perfect model for studying muscle cell signaling and therefore, our study encompassed rabbit and mouse muscle microsomes as well as a cultured mouse muscle cell

line. Most of what is known about DGC proteins is known from studies in rabbit muscle microsomes and this makes this an ideal system for the study of DGC protein interactions. However, cultured cells are most easily adapted to inhibitor studies and the *mdx* mouse is the best characterized animal model for Duchenne muscular dystrophy. Despite the number of model systems used, the results all show a high degree of internal consistency and give some confidence that the results reported are accurate for many skeletal muscle models.

Taken together, our results demonstrate that the DGC is associated with PI3K/Akt signaling and that DGC-G β γ is binding PI3K in a laminin-dependent manner and activating pAkt. These results are consistent across a variety of model systems for studying muscle signaling. Further downstream, this leads to GSK-3 β phosphorylation. *Mdx* mouse displays elevated integrin β 1 and increased interaction between integrin β 1 and laminin, G β γ and PI3K (Fig. 8) and increased Akt signaling compared to normal tissue. *Mdx* mice might compensate for the loss of dystrophin and associated DGC proteins by this enhance laminin-integrin β 1 interaction. Moreover, disruption of the DGC in *mdx* mouse is causing dis-regulation of the laminin-DGC-G β γ -PI3K-Akt signaling and replaces it with laminin-integrin-G β γ -PI3K-Akt signaling. This further substantiates the hypothesis that this laminin-DGC-G β γ -PI3K-Akt-GSK signaling is likely to be important to the pathogenesis of muscular dystrophies, and modulation of this pathway may be of therapeutic value for DMD. Most β 2-adrenergic agonists have the undesirable side-effect of inducing unwanted cardiac hypertrophy. However, some of the newer agonist, such as formoterol and salmeterol, also induce skeletal muscle hypertrophy with greater selectivity and by stimulating G β γ -induced pAkt activation may be of therapeutic value for DMD and other muscle wasting diseases (Lynch et al., 2007; Ryall et al., 2006).

Acknowledgments

We appreciate the generous gifts of antibodies from Drs. Tamara Petrucci and Kevin P. Campbell. The excellent technical assistance of Maria Macias and Magda Loranc is greatly appreciated. Yongmin Xiong is a visiting professor from Xi'an Jiaotong University School of Medicine; this work is in partial fulfillment of the requirements for her Ph.D.

Grants: This study was supported by National Institutes of Health grant AR051440, the Muscular Dystrophy Association grant #3789 and the China Scholarship Council.

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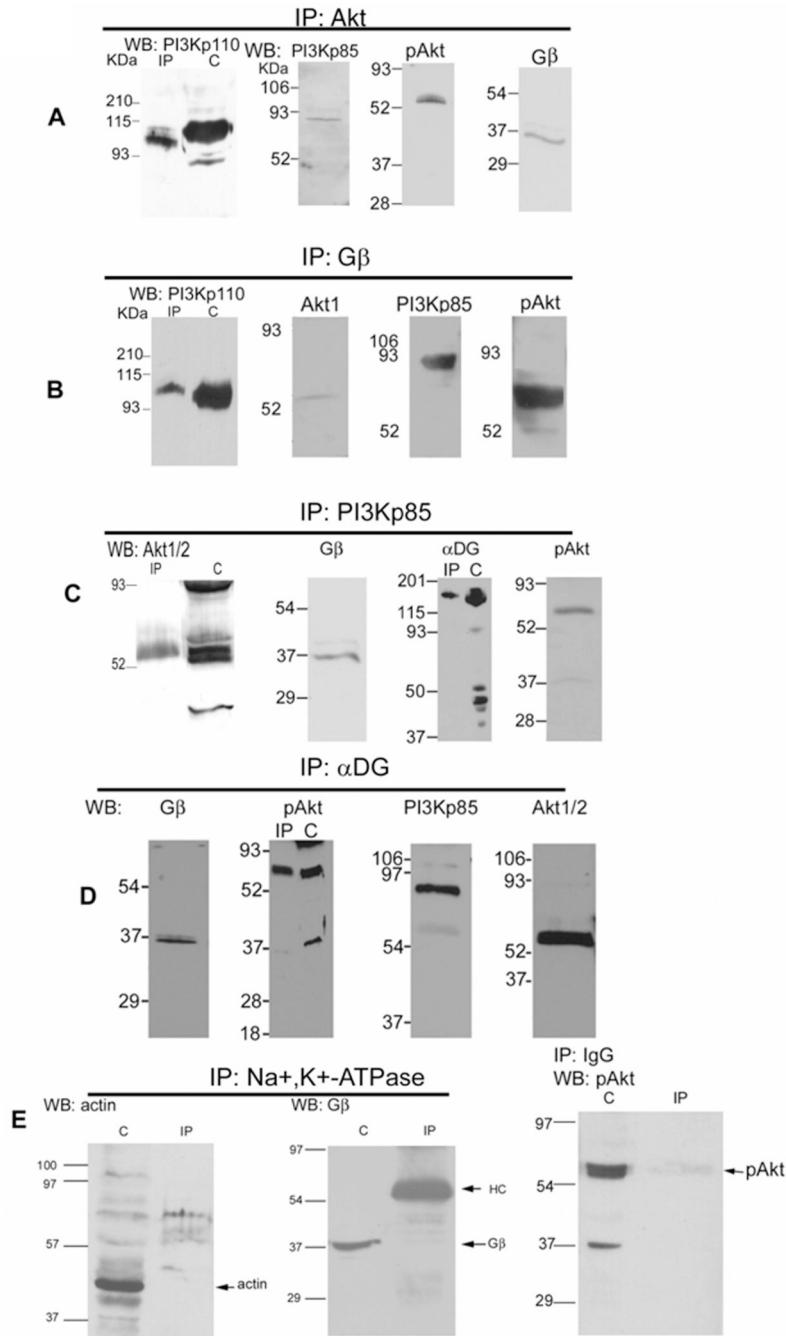


Fig. 1. DGC-Gβγ binds PI3K and Akt in rabbit skeletal muscle

A, Rabbit muscle microsomes were isolated and the protein concentration (Bradford, 1976) adjusted so that equal amounts were used for all experiments. Microsomes were incubated with antibodies cross-reactive with both Akt 1 and 2 (Akt 1/2), detergent solubilized upon addition of protein G-Sepharose, electrophoresed, electroblotted, and Western blotted for PI3Kp85, pAkt 1/2/3, PI3Kp110, and Gβ. Where shown “C” denotes crude microsomes prior to IP, added in equal concentration. IP: Immunoprecipitation, WB: Western blot. **B**, Microsomes were immunoprecipitated with a Gβ antibody, and Western blotted for PI3Kp110, PI3Kp85, Akt 1 and pAkt 1/2/3. **C**, Microsomes were immunoprecipitated with a PI3Kp85 antibody, and Western blotted for Akt1/2, α-dystroglycan (αDG, VIA4 monoclonal antibody), Gβ, and

pAkt1/2/3. **D**, Microsomes were immunoprecipitated with an α -dystroglycan monoclonal antibody (VIA4), and Western blotted for G β , PI3Kp85, pAkt and Akt1/2. These show that DGC-G $\beta\gamma$ binds a complex containing PI3K and Akt, as well as the phosphorylated Akt. **E**, Microsomes (C) or detergent immunoprecipitated with Na⁺/K⁺ ATPase or pre-immune IgG antibodies, were probed with the antibodies against actin, G β and pAkt.

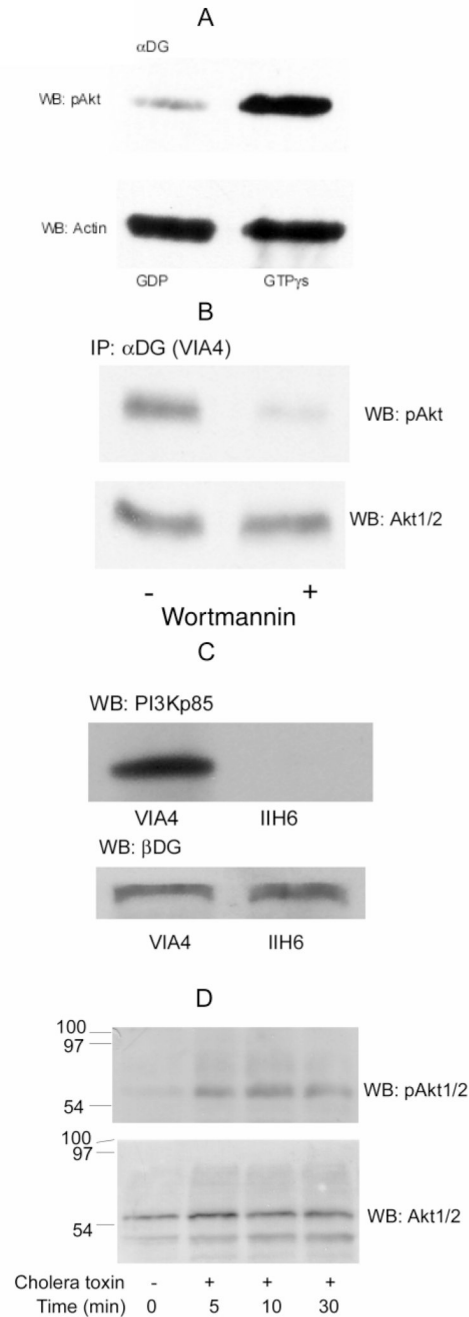


Fig. 2. Activation of PI3K/Akt by G β γ -dimers in skeletal muscles

A, Rabbit skeletal muscle microsomes were isolated and incubated for 1 h at 37° in buffer K containing 1 mM CaCl₂ and 1 mM ATP and either 1 mM GTP- γ S or 1 mM GDP, and then were immunoprecipitated with VIA4-1 α -dystroglycan (α -DG) antibody and Western blotted for pAkt. Membrane were then stripped and re-probed with an actin antibody, as indicated. pAkt was decreased in the presence of GDP compared to GTP- γ S, but actin was not significantly altered and provides a loading control. **B**, C2C12 myotubes were pretreated with 200 nM wortmannin overnight at 37°C (+) or with buffer alone (-). The myotubes were counted and adjusted to equal cell counts, microsomes prepared and immunoprecipitated with α DG antibody (VIA4). After SDS-PAGE and electroblotting, the blot was probed with antibody

against pAkt. The same samples loaded on adjacent wells of the same gel was probed for total Akt1/2 as a loading control. **C**, the myotube, grown in either antibody, were used to prepare microsomes, which were then pretreated with the same antibody against α DG (VIA4 or IIH6). After immunoprecipitation with protein G-Sepharose, the proteins were separated on SDS-PAGE and transferred to nitrocellulose membranes. The membrane was probed with antibody against PI3Kp85 (upper panel). The same blot was stripped and re-probed with a β DG antibody (lower panel) to confirm equal loading. **D**, An equal number of C2C12 cells were cultured with 100 ng/ml cholera toxin for 0, 5, 10, 30, 60 min at 37°. Clarified cell lysates were adjusted to the same protein concentration (Bradford, 1976) and equal amounts separated on 12% SDS-PAGE and transferred to nitrocellulose membranes. The membranes were probed with antibodies against pAkt (upper panel). The stripped blot from was re-probed for total Akt1/2 to confirm equal loading (lower panel).

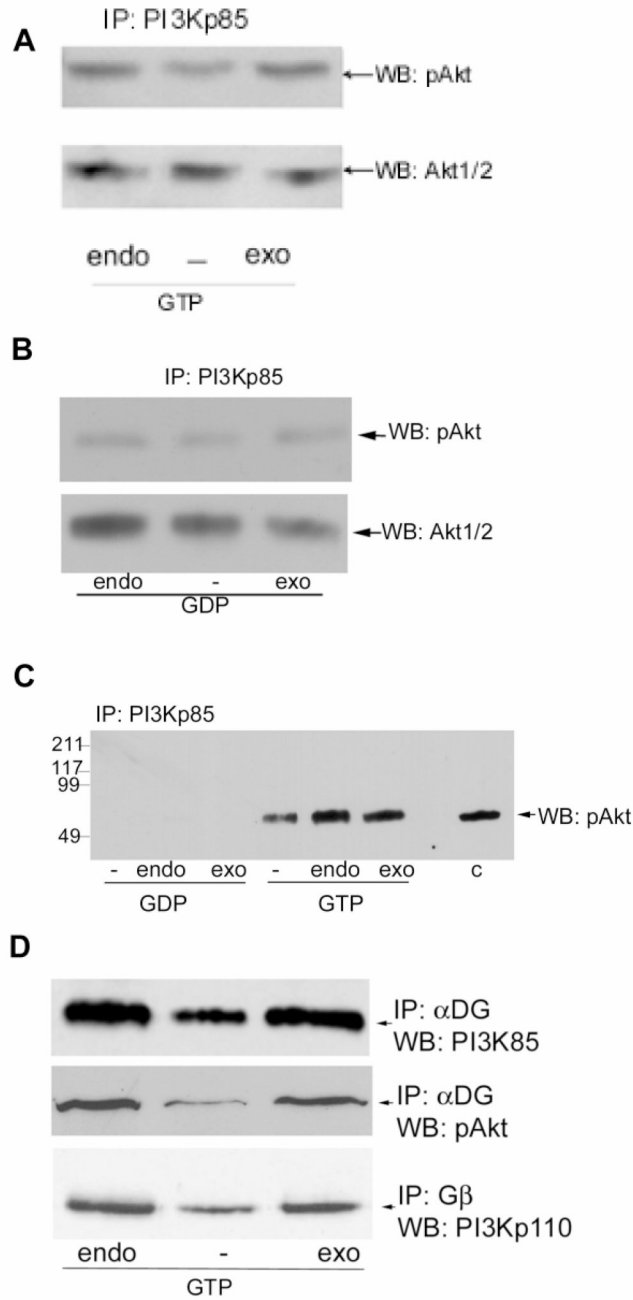


Fig. 3. Laminin binding alters PI3K/Akt signaling

A and **B**, rabbit skeletal muscle microsomes were treated with Sepharose 4B (endo) or with heparin-Sepharose to deplete laminin (-) for 1 hour at 4°. The microsomes were then adjusted to equal protein concentration (Bradford, 1976) and were made 1 mM CaCl₂, 10 mM MgCl₂, 1 mM ATP, and 1 mM GTP-γS (panel A) or 1 mM GDP (panel B). To the laminin-depleted portions of the microsome 3 μg exogenous laminin-1 (exo) was added for 1 hour at 4°, with gentle mixing. The microsomes were immunoprecipitated for PI3Kp85α and Western blotted with antibody against phosphorylated Akt (pAkt). Membrane were then stripped and re-probed with antibody to total Akt to show total (phosphorylated and unphosphorylated) protein levels in equal loads. The level of Akt phosphorylation was reduced in the laminin-depleted

immunoprecipitate, and laminin addition induces the phosphorylation of Akt. **C**, Microsomes were treated, incubated and immuno-precipitated as in panels A and B, except that the GTP- γ S and GDP treated samples and an unequal amount of untreated microsomes (C) were applied to the same gel, and Western blotted for pAkt on the same blot. The data show that much greater amounts of pAkt are immunoprecipitated from microsomes incubated with GTP- γ S than GDP and that laminin is also affecting the phosphorylation of pAkt.. **D**, Rabbit skeletal muscle microsomes were treated as in panel A. Immunoprecipitates with the α -dystroglycan (α -DG) VIA4 or G β antibody as indicated were probed with antibodies against PI3Kp85, PI3Kp110 or pAkt. Laminin binding increased α DG-G β -PI3K-pAkt interactions. Depletion of laminin resulted in decreased DGC-G β -PI3K-Akt interactions, and exogenous laminin-1 increases PI3K binding and activation of pAkt.

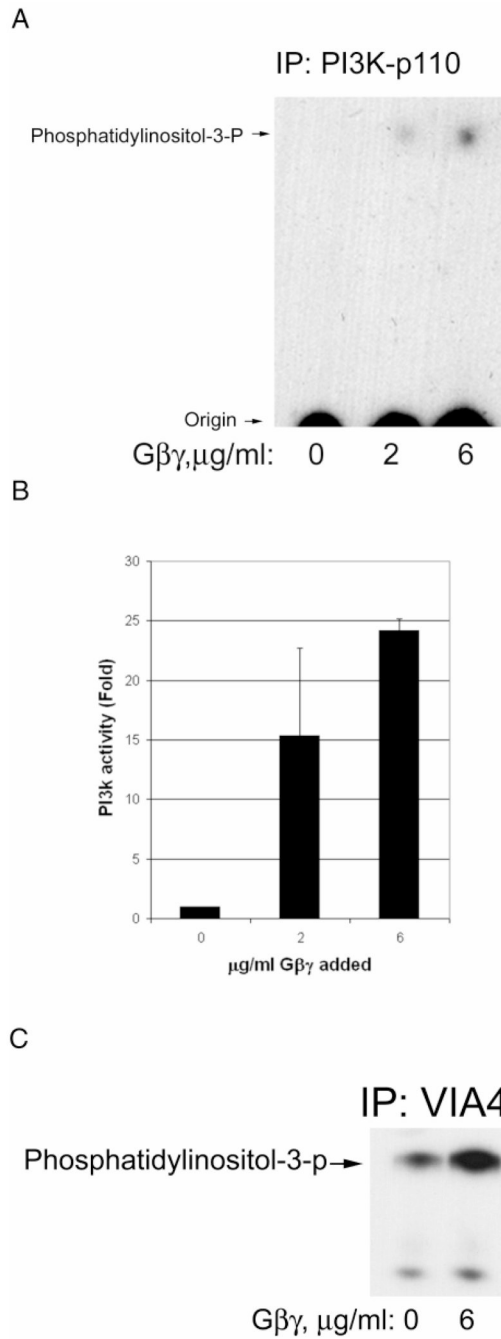


Fig. 4. Purified Gβγ activates PI3K, which is associated with the DGC

A, Rabbit skeletal muscle microsomes were incubated with GDP and the concentrations of purified Gβγ shown for 15 min. Then digitonin was added for solubilization followed by immunoprecipitation for PI3Kp110 catalytic subunit. The immunoprecipitate was added to a PI3K assay where phosphatidylinositol is added along with γ -³²P-ATP as described in Methods. The reaction mixture is extracted for lipids and analyzed by silica thin layer chromatography and autoradiography. The position of product phosphatidylinositol-3-phosphate, obtained from a standard also applied to the plate, is shown to the left. **B**, The same as panel A except that densitometry was performed and the data normalized so that the absence of Gβγ was one-fold activation. The bars represent the mean of two independent experiments

and the standard deviation is shown. C, The same as panel A except that VIA4 antibody was used to immunoprecipitate the DGC. Only the upper portion of the chromatogram is shown and at a longer exposure to emphasize an additional 3-phospholipid also detected.

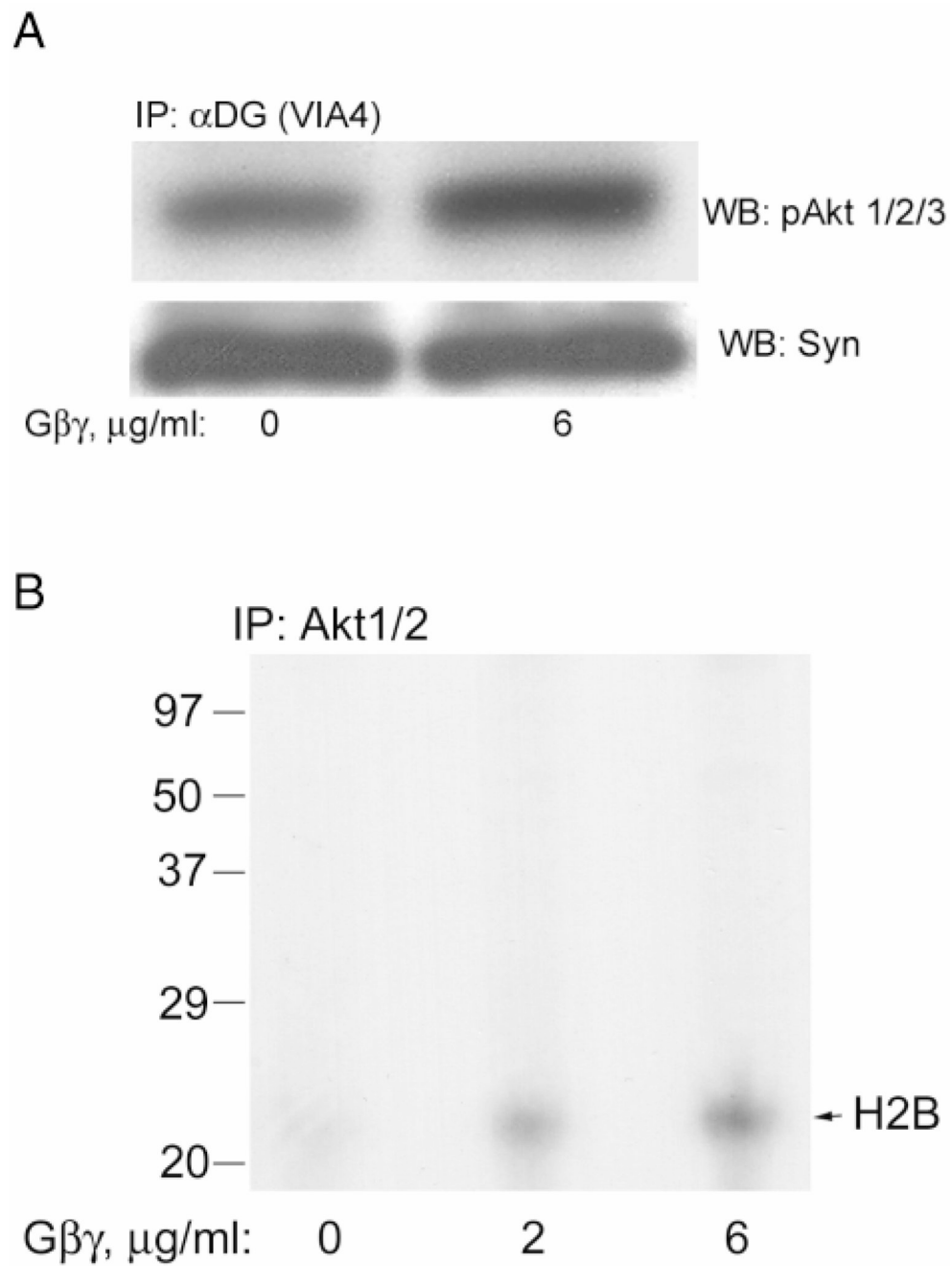


Fig. 5. Purified G $\beta\gamma$ activates Akt, which is associated with the DGC

A, Microsomes were treated as described in Fig. 4A and immunoprecipitated with VIA4 antibody and Western blotted. The pAkt antibody shows that pAkt was activated. The same blot was stripped and probed with the syntrophin antibody to provide a loading control. **B**, Immunoprecipitates were prepared as in panel A except with the Akt1/2 antibody and added to an Akt assay using histone as substrate. The autoradiogram is shown and the expected position of histone H2B is shown to the right.

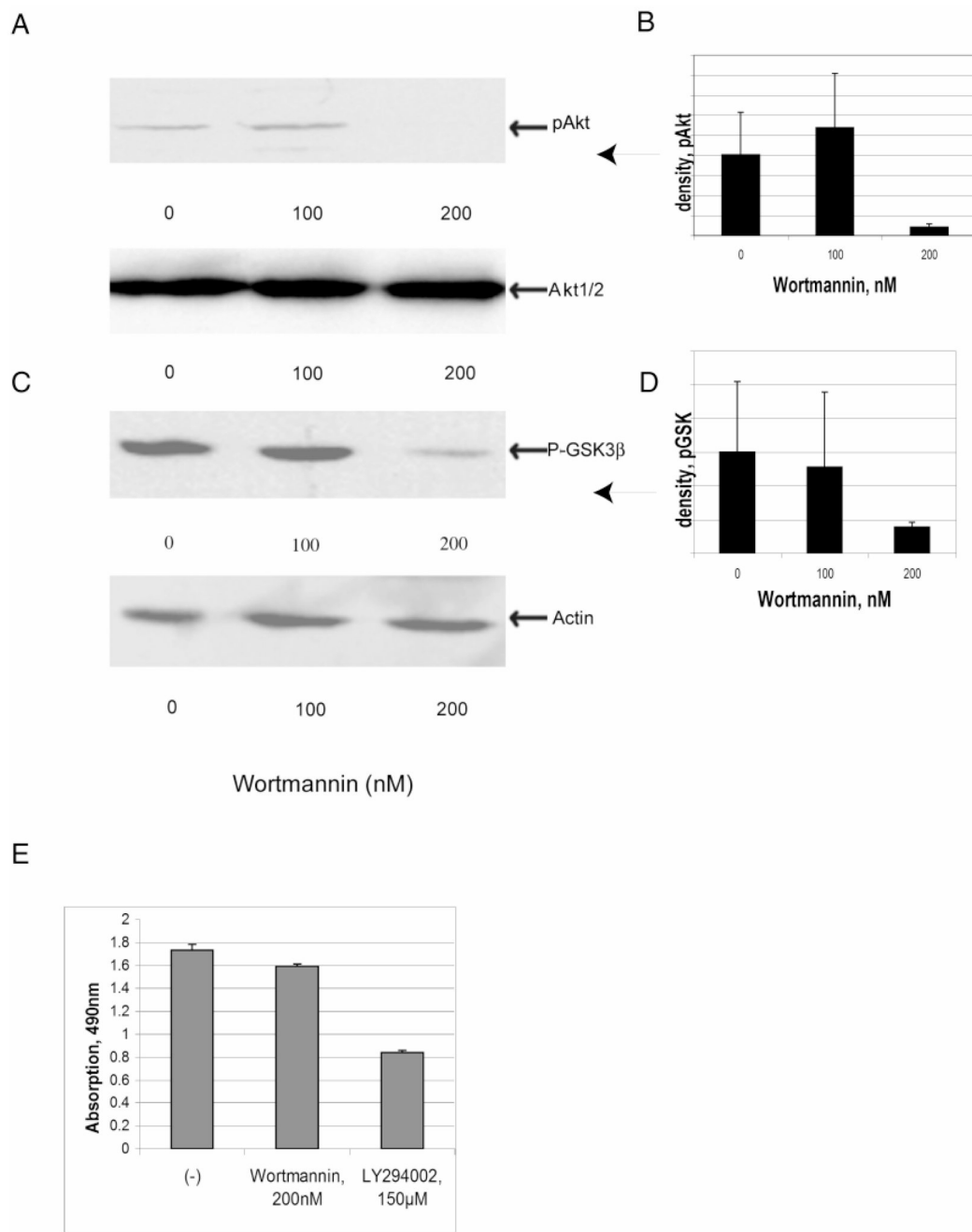
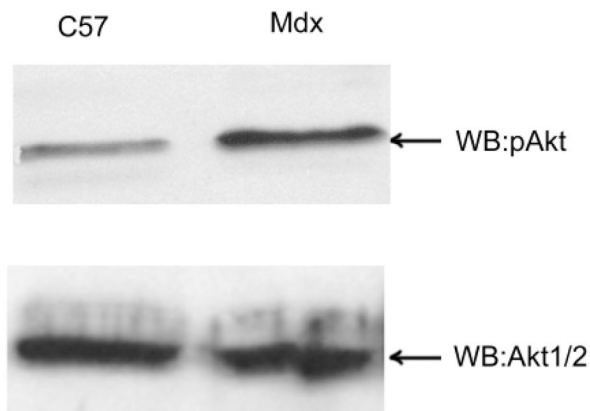


Fig. 6. Effects of Wortmannin on PI3K activation in C2C12 myotube cultures

Myotube cultures were maintained in 1% FBS DMEM medium for 6 days before addition of inhibitors. Myotubes were incubated in the absence or presence of different concentrations of Wortmannin as indicated in the figure. After 24 h, cells were lysed and subjected to Western blot assay. **A**, Western blots for pAkt are shown above with total Akt (on the same blot after stripping) below providing a loading control. **B**, Densitometry of the pAkt staining. **C**, Western blots for pGSK3β are shown above with actin shown below providing a loading control. **D**, Densitometry of the pGSK3 staining. **E**, The effect of Wortmannin and LY294002 on C2C12 myotube viability was measured using the MTS assay. The error bars throughout are standard deviation with n=2 for panels B and D, and n=3 for panel E.

A



B

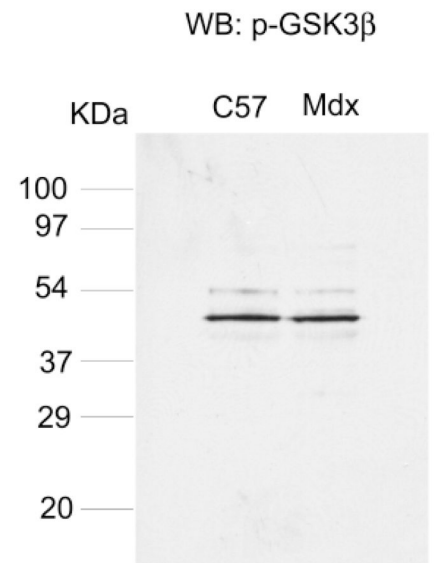


Fig. 7. Activation (phosphorylation) of Akt is increased in dystrophin-deficient skeletal muscles
A, Skeletal muscle microsomes from 4-week-old normal and *mdx* mice were prepared from the same wet weight of gastrocnemius muscle, separated on 12% SDS-PAGE and transferred to nitrocellulose. The blots were probed with antibody against pAkt, and membranes were then stripped and reprobed with the Akt 1/2 antibody to detect total Akt levels. pAkt (Ser 473) was elevated in 4-week-old muscle from *mdx* mice compared to muscle isolate from age-matched normal mice (C57) while total Akt was not. **B**, Skeletal muscle (gastrocnemius) microsomes from adult (6 month old) normal and *mdx* mice were probed for the phosphorylated, p-GSK3β, as indicated. pGSK was not different between normal and *mdx* mice.

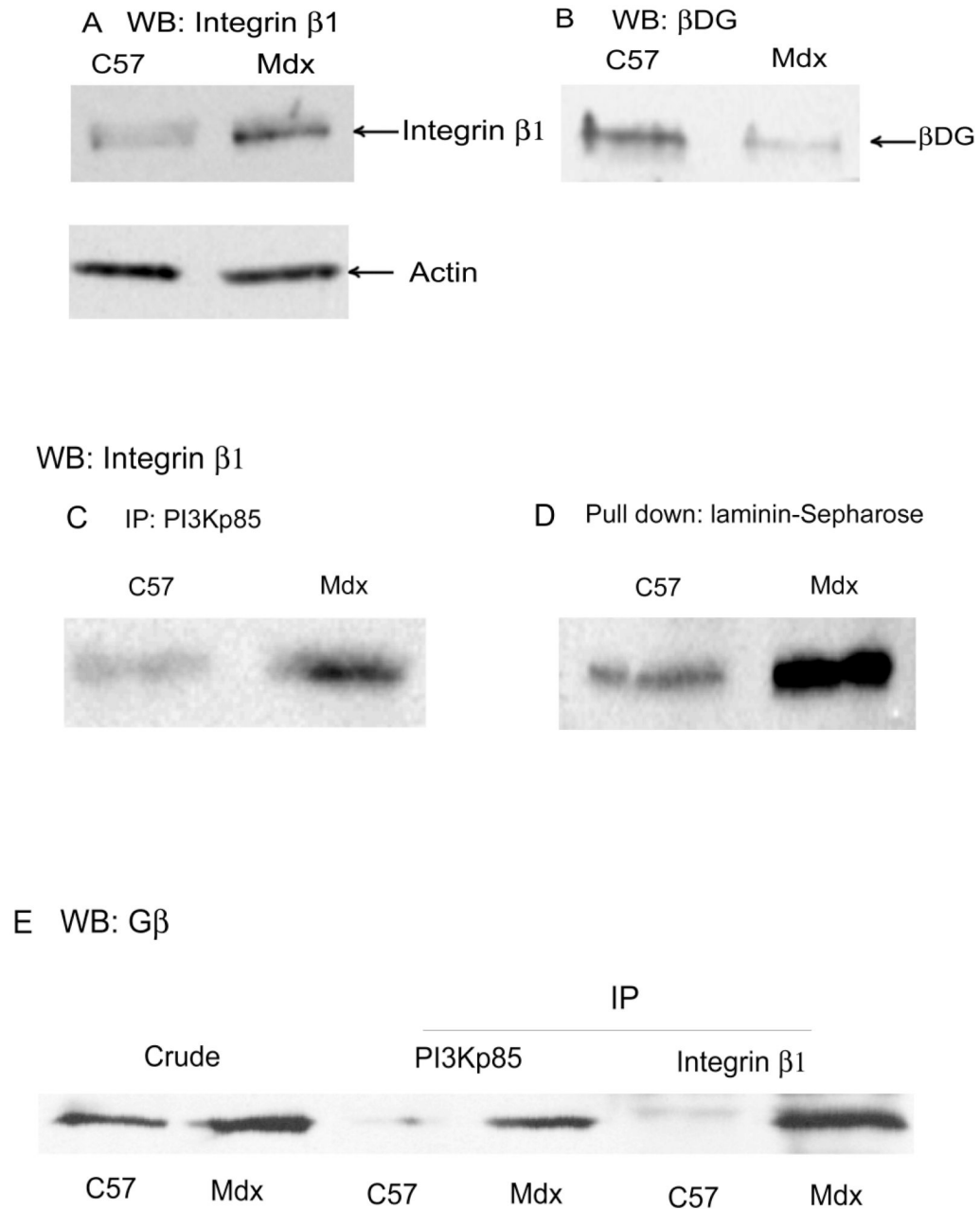


Fig. 8. Enhanced expression of integrin $\beta 1$ and G β in *mdx* muscle induces PI3K recruitment
A, Skeletal muscle (gastrocnemius) crude microsomes from adult (6 month old) normal (C57bl) and *mdx* mice, prepared from the same wet weight of muscle, were Western blotted with anti-integrin $\beta 1$ or (after stripping the same blot) anti-actin antibodies. Integrin $\beta 1$ protein expression was found to be higher in *mdx* mouse muscle compared with control while actin was not. **B**, Same as panel A except Western blotting with anti- βDG (shown) or anti-actin antibodies (data not shown). βDG was decreased in *mdx* muscle compared to muscle isolate from age-matched C57bl mice (shown) while actin was not significantly changed (data not shown). **C**, microsomes from 6 mo. old C57bl and *mdx* mice were immunoprecipitated with antibodies against PI3Kp85 and analyzed by Western blotting with antibody to integrin $\beta 1$.

Interaction of PI3K-integrin $\beta 1$ was enhanced in *mdx* muscle compared with normal muscle. **D**, Same as panel C except laminin-Sepharose was used instead of immunoprecipitation and again Western blotting with integrin $\beta 1$ antibody. Laminin-bound integrin $\beta 1$ increases in skeletal muscle of *mdx* compared to normal mice. **E**, Skeletal muscle microsomes (crude) from adult C57 and *mdx* mice and its immune precipitates (IP) with antibodies against PI3Kp85 and integrin $\beta 1$ were analyzed by Western blotting with antibody against G β . The results show that G β was increased in *mdx* muscle microsome compared with normal, and interactions of PI3K-G β and integrin $\beta 1$ -G β were enhanced in *mdx* mice.