

NIH Public Access

Author Manuscript

IUBMB Life. Author manuscript; available in PMC 2010 June 15

Published in final edited form as:

IUBMB Life. 2008 March ; 60(3): 145–153. doi:10.1002/iub.21.

Signaling Mechanisms in Skeletal Muscle: Acute Responses and Chronic Adaptations to Exercise

Katja S.C. Röckl, Carol A. Witczak, and Laurie J. Goodyear

Research Division, Joslin Diabetes Center and Department of Medicine, Brigham and Women's Hospital and Harvard Medical School, Boston, MA, USA

Summary

Physical activity elicits physiological responses in skeletal muscle that result in a number of health benefits, in particular in disease states, such as type 2 diabetes. An acute bout of exercise/muscle contraction improves glucose homeostasis by increasing skeletal muscle glucose uptake, while chronic exercise training induces alterations in the expression of metabolic genes, such as those involved in muscle fiber type, mitochondrial biogenesis, or glucose transporter 4 (GLUT4) protein levels. A primary goal of exercise research is to elucidate the mechanisms that regulate these important metabolic and transcriptional events in skeletal muscle. In this review, we briefly summarize the current literature describing the molecular signals underlying skeletal muscle signaling proteins involved in glucose transport, muscle fiber type, and mitochondrial biogenesis is ongoing. Further research is needed because full elucidation of exercise-mediated signaling pathways would represent a significant step toward the development of new pharmacological targets for the treatment of metabolic diseases such as type 2 diabetes.

Keywords

skeletal muscle; AMPK; glucose uptake; training adaptation; exercise

INTRODUCTION

Throughout the world, diabetes afflicts over 180 million people, and epidemiological estimates from the World Health Organization project that this number will reach 366 million (4.4% of the world population) by 2030. Thus, diabetes is rapidly being recognized as a public health threat that is rising to epidemic proportions. While the rate of diabetes is on the increase, it has long been recognized that physical activity has important health benefits for people with type 2 diabetes. In the acute state, exercise positively moderates glucose homeostasis by enhancing glucose transport and insulin action in contracting skeletal muscle, the major tissue responsible for total body glucose disposal (1). Chronic physical activity (i.e. exercise training) increases glucose transporter 4 (GLUT4) protein levels, mitochondrial enzyme content, and alters fiber type in skeletal muscle (2,3), thus providing an additional mechanism for exercise-mediated improvements in insulin sensitivity. Collectively, these effects help to explain the strong epidemiological evidence that regular physical activity prevents or delays the onset of type 2 diabetes (4,5).

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Address correspondence to: Laurie J. Goodyear, Senior Investigator and Head, Metabolism Section, Joslin Diabetes Center, One Joslin Place, Boston, MA 02215, USA. Tel: + 617-732-2573; Fax: 617-732-2650. laurie.goodyear@joslin.harvard.edu.

Despite the physiological importance of exercise in regulating skeletal muscle metabolism, the molecular mechanisms that underlie these important phenomena are only partly understood. Elucidating exercise-stimulated and insulin-independent signals that mediate glucose transport have already led to the identification of new targets for the treatment of diabetes (*e.g.*, AMP-activated protein kinase (AMPK)). Determining the comprehensive mechanism will undoubtedly provide more targets for treatment, as well as provide fundamental knowledge of this complex physiological process. In this manuscript, we will briefly review the current literature on this important area of exercise and diabetes research.

ACUTE EFFECTS OF EXERCISE: REGULATION OF SKELETAL MUSCLE GLUCOSE TRANSPORT

Insulin and exercise are the two most physiologically relevant stimulators of skeletal muscle glucose transport (6,7). In individuals with type 2 diabetes, the insulin-dependent regulation of skeletal muscle glucose transport is impaired. Importantly, insulin independent mechanisms, including exercise/contraction-mediated mechanisms for regulating glucose uptake remain intact (8,9). Both insulin and exercise/muscle contraction increase skeletal muscle glucose uptake by translocation of glucose transporters from an intracellular location to the plasma membrane and t-tubules. GLUT4 is the predominant glucose transporter isoform expressed in skeletal muscle. Our laboratory and others have worked to elucidate the signaling mechanisms leading to exercise-stimulated GLUT4 translocation (6,7). Early studies have demonstrated that there are distinct proximal signaling mechanisms responsible for the stimulation of GLUT4 translocation and glucose transport by insulin and exercise. Insulin signaling involves the rapid phosphorylation of the insulin receptor, insulin receptor substrate-1/2 (IRS-1/2) on tyrosine residues, and the activation of phosphatidylinositol 3-kinase (PI3-K) (10,11). In contrast, exercise and muscle contraction have no effect on insulin receptor and IRS-1 phosphorylation or on PI3-K activity (11,12), and muscle-specific knockout of the insulin receptor does not impair contraction-stimulated glucose transport (13). Clearly, these data demonstrate that the initiating signals that lead to GLUT4 translocation by insulin and exercise in skeletal muscle are distinct.

Muscle contraction is a multifactorial process involving changes in cellular energy status (i.e. increased AMP:ATP), increases in intracellular Ca^{2+} levels, activation of protein kinase C (PKC), and so forth. Not surprisingly, this has led many investigators to speculate that these processes activate one or more intracellular signaling pathways that coordinately act to increase plasma membrane GLUT4 transporters and glucose uptake in response to physical activity. In the following sections, we will discuss the signaling proteins that have been implicated in this complex process, including AMPK, LKB1, Ca^{2+} /calmodulin-dependent protein kinases (CaMKs), PKCs, and AS160 (Fig. 1).

AMP-activated Protein Kinase

The AMPK and its primary upstream kinase, LKB1, are the most widely studied proteins implicated in muscle glucose transport in response to changes in the cellular energy status. AMPK is a heterotrimeric protein that is activated by a complex mechanism involving an increase in the AMP:ATP ratio and allosteric modification and phosphorylation by one or more upstream kinases, including LKB1 (14,15). Studies using the AMP-analog, 5-aminoimidazole-4-carboxamide ribonucleoside (AICAR), have demonstrated that activation of AMPK is positively correlated with an increase in muscle glucose uptake (16). However, data obtained from mouse models of attenuated AMPK activity demonstrate that inhibition of AMPK has little or no effect on contraction-induced glucose uptake (17–19). Furthermore, muscle-specific ablation of LKB1 only partially inhibits contraction-stimulated glucose

transport (20,21). Collectively, these data suggest that various signals are involved in the regulation of contraction-stimulated glucose transport in skeletal muscle.

Ca²⁺/Calmodulin-dependent Protein Kinases

Increases in intracellular Ca²⁺ levels are a fundamental part of muscle contraction, and recent studies have implicated Ca²⁺/calmodulin signaling and Ca²⁺/calmodulin-dependent protein kinases as critical molecules underlying Ca²⁺ - and contraction-stimulated skeletal muscle glucose transport. Incubation of rat epitrochlearis muscles with the Ca²⁺/calmodulin competitive inhibitor, KN-93, decreases skeletal muscle glucose transport in response to contraction and to stimulation with the sarcoplasmic reticulum Ca²⁺ store releasing agent, caffeine (22). In the same study, KN-93 significantly inhibited contraction-induced CaMKII phosphorylation in the absence of AMPK inhibition, suggesting that CaMKs regulate glucose uptake independent of AMPK signaling. These results are consistent with a recent study from our lab, demonstrating that CaMK kinase α -dependent signaling can stimulate muscle glucose uptake without changes in AMPK activity (23). In contrast, a recent study using isolated mouse muscles has shown that inhibition of CaMK signaling with KN-93 or the CaMK kinase inhibitor, STO-609, inhibits contraction-induced skeletal muscle glucose uptake via an AMPK-dependent signaling pathway (24). Thus, the role of AMPK in the regulation of Ca²⁺/caMK kinase in skeletal muscle glucose uptake is still being debated.

Protein Kinase C

PKC is another molecule which is activated by muscle contraction and has been implicated in the regulation of contraction-stimulated muscle glucose transport (25,26). In mammalian cells, 12 different PKC isoforms have been identified and classified into three subfamilies based on amino acid similarity and mode of activation (27). The conventional PKCs (cPKCs, α , β 1, β 2, and γ isoforms) are dependent on Ca²⁺ and diacyl-glycerol for activation, the novel PKCs (nPKCs, δ , ε , θ , and η isoforms) are dependent on only diacylglycerol for activation, and the atypical PKCs (aPKCs, ζ and λ isoforms) are activated independently of both Ca²⁺ and diacylglycerol. Pharmacological inhibition of cPKCs and nPKCs blunts contraction-stimulated skeletal muscle glucose uptake in skeletal muscle (25,26), suggesting that PKCs are important in this process. However, recent studies assessing isoform-specific PKC activation have failed to demonstrate an increase in cPKC or nPKC activity by exercise/contraction (28-30). This controversy might be explained by a certain degree of nonspecificity of the pharmacological compounds. In contrast to the findings on cPKCs and nPKCs, our laboratory and others have recently shown that aPKCs are activated by exercise (29,31,32). In addition, in 3T3-L1 adipocytes and L6 myotubes, overexpression of wild type and constitutively active aPKCs stimulates or potentiates the effects of insulin on glucose transport, while kinase-inactive aPKCs inhibit the effect of insulin on glucose transport (33-35). Other data in support of a role for aPKCs in the regulation of glucose transport comes from a study where PKC ζ and λ were hypothesized to be downstream of AMPK (31). This study, performed in L6 myotubes and isolated rat muscles, suggests that the effects of AMPK on glucose transport are mediated through the sequential activation of the extracellular signal-regulated kinase (ERK), prolinerich tyrosine kinase-2, phospholipase D, and aPKCs (31). These results are not entirely consistent with our previous data which show insulin- and contraction-stimulated glucose transport to be independent of ERK signaling (36,37). In future studies, it will be important to further elucidate a putative AMPK-aPKC interaction.

Akt Substrate of 160 kDa (AS160)

The Akt substrate of 160 kDa (AS160) is a recently discovered protein that regulates insulinstimulated GLUT4 translocation in 3T3-L1 adipocytes (38), L6 myotubes (39), and rat skeletal muscle (40). It is phosphorylated on six different phospho-Akt-substrate (PAS) sites in

response to both insulin and contraction in skeletal muscle (40–42). Recent evidence from our lab has shown that AMPK phosphorylates AS160 (PAS) in response to AICAR and contraction in skeletal muscle (41), and that mutation of four PAS sites significantly inhibits both insulin and contraction-induced glucose uptake (43). In addition, exciting new data from our lab demonstrates that mutation of the calmodulin-binding domain on AS160 significantly inhibits contraction- but not insulin-stimulated glucose uptake (44). Thus, both phosphorylation and calmodulin-binding on AS160 appear to play a role in the regulation of contraction-stimulated glucose uptake. Collectively, these results suggest that AS160 may serve as a point of convergence for both insulin- and contraction-dependent signaling in the regulation of glucose uptake.

Other Putative Mediators of Skeletal Muscle Glucose Uptake

Investigators have speculated on the possible involvement of other signals in the regulation of contraction-stimulated skeletal muscle glucose transport, including bradykinin, reactive oxygen species, and nitric oxide. Although, there is currently no evidence in support of a role for bradykinin in muscle glucose transport (45), several studies have provided data supporting a role for reactive oxygen species (46) and nitric oxide (47–50). However, further investigations have suggested that nitric oxide stimulates glucose transport *via* a mechanism distinct from contraction and insulin-dependent signaling pathways (48,49,51).

ADAPTATIONS OF SKELETAL MUSCLE TO EXERCISE TRAINING

Regular physical activity leads to a number of adaptations in skeletal muscle that allow the muscle to more efficiently utilize substrates for ATP production and thus become more resistant to fatigue. In the following sections, we will summarize the current literature on three major adaptations to exercise training: 1) muscle fiber type transformations as defined by the expression of specific contractile proteins (myosin heavy chain isoforms), 2) increases in mitochondrial activity and content, and 3) increases in GLUT4 protein expression.

Muscle fiber types have traditionally been classified according to their expression of myosin heavy chain isoforms as fast twitch fibers (type IIb, IIx, and IIa) and slow twitch fibers (type I). Type IIb and type IIx fibers mainly depend on glycolytic, and type IIa and type I fibers on oxidative pathways for ATP production (52–54). There is an association between fiber type and mitochondrial content, with type IIb fibers tending to have the lowest and type I fibers the highest abundance of mitochondria. Endurance exercise has been shown to induce an increase in mitochondrial content and activity within the same fiber type but also a change in myosin heavy chain isoform expression, thus provoking a fiber type transformation from type IIb to IIx and IIa, and in rare cases also to type I muscle fibers (54). It is important to understand that mitochondrial biogenesis and fiber type transformation can occur independent from each other, suggesting distinct signaling mechanisms for both types of adaptive responses. In addition to increases in slow muscle fiber types and mitochondria, GLUT4 protein expression also increases in response to exercise training, facilitating glucose uptake into the trained muscle (6). GLUT4 protein is expressed to the highest level in slow, oxidative fiber types (55). Thus, muscle fiber type, mitochondrial content, and the abundance of GLUT4 are associated within the muscle cell but appear to be regulated independently. Interestingly, individuals with insulin resistance or type 2 diabetes have a distinct muscle phenotype with decreased slow, oxidative muscle fibers (56,57), and decreased GLUT4 expression within the slow muscle fibers (58). Skeletal muscle mitochondrial dysfunction has also been linked with insulin resistance and type 2 diabetes (59), although more studies are needed to directly connect mitochondrial deficits with impaired muscle glucose metabolism. Taken together, these findings imply that the discussed aspects of muscle fiber plasticity play an important role in the pathogenesis of diabetes, and that the benefits of exercise training for people with diabetes may stem from the aforementioned training-induced skeletal muscle adaptations. In the following sections, we

will give a brief overview of the signaling molecules which are considered to play a key role in muscle fiber type transformation, mitochondrial biogenesis, and increased GLUT4 expression in response to exercise training (Fig. 2).

AMPK

The AMPK has been implicated in the regulation of muscle fiber type (60–62), mitochondrial biogenesis (63), and GLUT4 biogenesis in skeletal muscle (64), making AMPK a key protein of interest in the study of exercise-mediated muscle adaptations. In young rats, administration of the AMPK activator, AICAR, for 14 days (1 mg/g body wt, subcutaneous) significantly increased the number of type IIx fibers in extensor digitorum longus (61). In addition, recent work from our lab using AMPK γ 1 (R70Q) transgenic mice, a mouse model of constitutively active AMPK activity (65), demonstrated that chronic elevations in AMPK activity resulted in significant increases in the ratio of type IIa/x fibers in triceps muscles (62). Collectively, these results suggest a role for AMPK in exercise-induced muscle fiber type transformation. However, exercise training of AMPK α 2 (D157A) transgenic mice, mice with chronic inhibition AMPK α 2 activity (17), only showed a partial impairment in the exercise-induced increase in type IIa/x fibers (62). Thus, in addition to AMPK signaling, other pathways must be involved in training-induced muscle fiber type transformations.

Activation of AMPK with the AMP-analog AICAR or the creatine analog, β guanidinopropionic acid, increases the activity and content of mitochondrial proteins (63,66, 67), and this effect is abolished in AMPK α 2 kinase dead (67) and knockout mice (68). Consistent with these findings, transgenic mice with increased AMPK activity show increases in mitochondrial markers (62). Furthermore, chronic AICAR injections induce peroxisomeproliferator-activated receptor- γ coactivator 1 α (PGC-1 α) RNA expression in rat skeletal muscle (61,69), suggesting a possible link between AMPK activation and PGC-1 α signaling. Surprisingly, in response to exercise training, AMPK α 2 knockout and AMPK α 2 kinase inactive mice show no defect in mitochondrial increases (62,68). Thus, despite its capacity to induce mitochondrial biogenesis, AMPK is not essential for exercise training-induced increases in mitochondria.

Additional studies using AICAR demonstrated that chronic activation of AMPK (5 days) increased muscle GLUT4 expression ~50–200% (64), suggesting that AMPK may regulate muscle GLUT4 biogenesis. However in two recent studies, AMPKγ1 (R70Q) transgenic mice that exhibit chronic increases in AMPK activity, did not have increased muscle GLUT4 mRNA and protein expression (62,65). Thus, the involvement of AMPK in AICAR- or exercise-stimulated increases in GLUT4 protein levels is currently controversial.

Peroxisome-proliferator-activated Receptor-y Coactivator 1a

PGC-1 α is a potent transcriptional coactivator that interacts with a variety of transcription factors (*e.g.* MEF2, ERR α , NRF-1, NRF-2) to regulate glucose and fatty acid metabolism, mitochondrial biogenesis, and muscle fiber type transformation from type II to type I fibers (70–74). Both short-term exercise and endurance training stimulate PGC-1 α expression in myocytes (75), and this is thought to occur *via* a positive feedback mechanism involving increased expression of MEF2. In addition, in studies using *in vivo* gene transfection, mutations of the PGC-1 α promoter at MEF2 binding sites or a cAMP response element (CRE) show that contraction-induced PGC-1 α promoter activity in skeletal muscle is dependent on its MEF2 and CRE sequence elements (76). Research over the last several years has attempted to identify upstream signaling molecules involved in the activation of PGC-1 α including AMPK, the Ca²⁺/calmodulin-dependent protein phosphatase calcineurin, CaMKs, and p38 mitogen activated protein kinase (p38 MAPK) (69,73,77,78). In a recent study, AMPK was shown to induce increases in the binding of PGC-1 α to its promoter, via direct phosphorylation of

PGC-1 α on amino acids Thr¹⁷⁷ and Ser⁵³⁸ (79). Thus, a number of signaling pathways activate the transcriptional coactivator PGC-1 α to regulate muscle fiber types, mitochondria, and glucose metabolism. This suggests a pivotal role of PGC-1 α in skeletal muscle adaptations to exercise.

Calcineurin

The Ca²⁺/calmodulin-dependent protein phosphatase, calcineurin, is known as the master regulator of fast to slow twitch fiber type changes (80–82). In transgenic mice expressing a constitutively active form of calcineurin, there is a substantial increase in the number of slow twitch type I muscle fibers (83). Conversely, inhibition of calcineurin activity by treatment with cyclosporin (5 mg/kg, daily for 6 weeks) promotes slow-to-fast fiber type transformation (80). In C2C12 myocytes, calcineurin significantly increases the ability of PGC-1 α to activate slow fiber promoters, suggesting a cooperation between the calcineurin pathway and PGC-1 α (73).

Reports on the role of calcineurin in mitochondrial biogenesis have been controversial (84). Transgenic mice expressing constitutively active calcineurin show increased PGC-1 expression (85), and in cultured cardiac myocytes, constitutively active calcineurin upregulates a large number of genes involved in mitochondrial energy metabolism (86). Furthermore, transplant patients maintained on the calcineurin inhibitor cyclosporin A develop a loss of skeletal muscle oxidative capacity (87). Despite these data suggesting an important role of calcineurin in mitochondrial biogenesis, other studies have shown that the phosphatase cannot fully account for exercise training-induced adaptations in skeletal muscle. Cyclosporin treatment (500 ng/ml for 14 days) does not prevent the upregulation of mitochondrial markers in Ca²⁺ ionophore-treated myotubes (88)or in trained rats (20 mg/kg for 7 days (89) or 50 mg/kg for 10 days (90)).

A role for Ca²⁺-dependent signaling in skeletal muscle GLUT4 expression first emerged from experiments in L6 cells using the sarcoplasmic reticulum Ca²⁺ mobilizing agent, caffeine (91). In this study, intermittent caffeine treatment (5 mM, 3 h/day, 5 days) induced increases in cytosolic Ca²⁺ levels, and significantly increased GLUT4 protein levels (~50%), suggesting that Ca²⁺ signaling regulates muscle GLUT4 expression. In addition, transgenic mice overexpressing a constitutively active form of calcineurin have increased skeletal muscle GLUT4 protein, suggesting that calcineurin can induce GLUT4 biogenesis (85). However, cyclosporin treatment of rats (20 mg/kg, daily for 4 weeks) did not impair exercise training-induced increases in GLUT4 protein and mRNA expression despite complete inhibition of calcineurin (92). Thus, a physiological role of calcineurin in GLUT4 upregulation following exercise training is still being debated.

Ca²⁺/Calmodulin-dependent Protein Kinases

Similar to the findings from constitutively active calcineurin transgenic mice, transgenic mice expressing a constitutively active form of CaMKIV exhibit an abundance of slow twitch type I muscle fibers (77). However, the physiological relevance of these findings is questionable, since recent findings have established that CaMKIV protein is not expressed in mouse skeletal muscle (93). Future studies will be needed to investigate the potential relevance of other CaMK family members in skeletal muscle fiber type adaptations to exercise.

A role for CaMKs in muscle mitochondrial biogenesis first emerged from studies in L6 cells (a rat muscle cell culture model) using the Ca²⁺ ionophores, A-23187 (94) and ionomycin (95), and the sarcoplasmic reticulum Ca²⁺ mobilizing agent, caffeine (95,96). These studies demonstrated that intermittent or sustained increases in cytosolic Ca²⁺ levels significantly increased mitochondrial enzymes, including COX-I, δ -aminole-vulinate (ALAS), citrate

synthase, and cytochrome c, and that treatment with the Ca $^{2+}$ /calmodulin competitive inhibitor, KN-93 (10 μ M) completely blocked caffeine-induced increases in ALAS, COXI, cytochrome c, and citrate synthase protein expression. In addition, transgenic mice expressing a constitutively active form of CaMKIV exhibited significant increases in muscle mitochondrial mass, along with increases in cytochrome b, CPT-1, and PGC-1 expression (77). However, whole body CaMKIV knockout mice do not exhibit impaired muscle mitochondrial biogenesis in response to contractile activity, and, as mentioned earlier, CaMKIV protein is actually not detectable in mouse skeletal muscle (93). Thus, the potential significance of the findings in CaMKIV transgenic mice lies in a possible homology with other members of the CaMK family that are expressed in skeletal muscle.

As described earlier, increased cytosolic Ca^{2+} levels by caffeine treatment induce the expression of GLUT4 protein (91). In addition, pretreatment of cells with the Ca^{2+} /calmodulin competitive inhibitor, KN-93, completely prevents the caffeine-induced increase in GLUT4 protein. These findings implicate an important role of CaMKs in elevated GLUT4 protein expression following increases in intracellular Ca^{2+} .

p38 MAPK

The p38 mitogen activated protein kinase (p38 MAPK) is activated by various stimuli including contraction, insulin, environmental stress, and proinflammatory cytokines (97–99). p38 MAPK has been suggested to play a functional role in myogenic cell differentiation (100), and our lab and others have shown increased p38 MAPK activation following various muscle contraction or running exercise protocols in rodents and humans (101,102). Studies in C2C12 myocytes show p38 MAPK as an activator of the PGC-1 α promoter, and this activation is mediated by the transcription factor ATF2 (78). In transgenic mice, muscle specific activation of p38 MAPK leads to enhanced PGC-1 α gene expression and increased mitochondrial proteins (78). An acute bout of exercise in mice (3 h of voluntary wheel running) or rats (2 h of swimming) increases p38 MAPK and ATF2 phosphorylation, leading to PGC-1 α activation (78,103). Since studies have so far focused on acute effects, it will be an interesting area of future research to determine the role of p38 MAPK in exercise training-induced adaptations in skeletal muscle.

SUMMARY

Exercise is of critical importance for people with insulin resistance or diabetes. Our current understanding is that one of the many benefits of an acute bout of exercise is an insulinindependent increase in the glucose uptake capacities of skeletal muscle. Important chronic adaptations to exercise training are the increase of mitochondria and thus oxidative capacities in skeletal muscle, the transformation of muscle fiber types, and the increase in GLUT4 protein expression.

Contractile activity and insulin are the most potent and physiologically relevant stimuli of glucose transport in skeletal muscle. While significant progress has been made in elucidating the insulin signaling pathway leading to GLUT4 translocation, identification of the signals mediating contraction-stimulated glucose transport has proved challenging. A growing body of data suggests that multiple signaling cascades mediate the metabolic effects of contraction. While the proximal signals leading to contraction- and insulin-stimulated glucose transport are clearly distinct, emerging studies have shown a reconnection or convergence of these signals at AS160.

Exercise training induces an increase of oxidative capacity, fiber type changes, and elevated GLUT4 protein levels in skeletal muscle; adaptations which are of critical importance to lower free fatty acids, improve glucose uptake, and decrease the risk of insulin resistance and diabetes. Again, multiple signaling pathways appear to act synergistically to mediate adaptive responses

to exercise training. In particular, AMPK and calcineurin have evolved as major candidates for mediating exercise-training adaptations. PGC-1 α may be a point of convergence for both pathways. While considerable progress has been made in decoding molecular mechanisms around these molecules, more research will be needed to test their physiological role in skeletal muscle adaptations to exercise training.

Acknowledgments

Work in the authors' laboratory was supported by grants to L.J. Goodyear (National Institutes of Health R01AR45670 and R01DK068626) and a Diabetes Endocrinology Research Grant at the Joslin Diabetes Center (National Institutes of Health DK36836). C.A. Witczak was supported by fellowships from the National Institutes of Health (F32AR051663) and the American Diabetes Association (mentor-based to L.J.G.). K.S.C. Röckl was supported by a fellowship within the Postdoc-Program of the German Academic Exchange Service, DAAD.

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Figure 1.

Proposed model for the signaling pathways mediating insulin and contraction-induced skeletal muscle glucose transport. Insulin and contraction-mediated glucose transport occurs by translocation of glucose transporter 4 (GLUT4) from intracellular locations to the plasma membrane. Insulin binding leads to phosphorylation of the insulin receptor with subsequent activation of insulin receptor substrate 1/2 (IRS-1/2) and phosphatidylinositol 3-kinase (PI3-kinase). Downstream of PI3-kinase the protein kinases, Akt, which then regulates activation of Akt Substrate of 160 kD (AS160), and atypical protein kinase C (aPKC), have been identified to mediate insulin stimulated GLUT4 translocation. Contraction stimulated glucose uptake is mediated by multiple signaling pathways including aPKC, Ca^{2+} /calmodulin-dependent protein kinase (I (CaMKII), Ca^{2+} /calmodulin-dependent protein kinase kinase (CaMKK), LKB1, and AMP-activated protein kinase (AMPK).



Figure 2.

Proposed model for the signaling pathways mediating fiber type transformation, mitochondrial biogenesis, and GLUT4 protein expression with skeletal muscle adaptations to endurance training. Exercise training leads to skeletal muscle fiber type transformation, mitochondrial biogenesis, and increased glucose transporter 4 (GLUT4) protein expression, and multiple signaling pathways have been suggested to be involved in these adaptations. Changes in the cellular energy status (AMP:ATP) stimulate AMP-activated protein kinase in the presence of the AMPK kinase, LKB1. AMPK may be involved in fiber type transformation, mitochondrial biogenesis, and GLUT4 biogenesis through increasing peroxisome-proliferator-activated receptor- γ coactivator 1 α (PGC-1 α) expression and probably also independent of PGC-1 α . Exercise training-induced increases in PGC-1 α are potentiated by a positive feedback loop through myocyte-enhancing factor 2 (MEF2) and are involved in fiber type transformation, mitochondrial biogenesis, and increased GLUT4 expression. Increases in intracellular Ca²⁺ levels lead to activation of the Ca²⁺/calmodulin-dependent phosphatase, calcineurin, as well as Ca²⁺/calmodulin-dependent protein kinases (CaMKs). While calcineurin is involved in a number of skeletal muscle adaptations, acting primarily through PGC-1 α , a role of CaMKs has so far pointed toward increasing GLUT4 protein expression. Contraction-induced activation of p38 mitogen activated protein kinase (p38 MAPK) increases PGC-1α expression through activating transcription factor 2 (ATF2) and may therefore also be involved in skeletal muscle adaptations to exercise training.