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## Chapter 5. Physiological and molecular adaptations to strength training

**Abstract.** High muscle contraction forces that lead to gains in muscle function, size, and strength characterize resistance exercise training. The purpose of this chapter is outline the adaptations in myofiber size and metabolism that occur by stimuli of hormones and local growth factors, mechanical and metabolic stress of muscle tissue, and myofibrillar disruptions induced by a resistance exercise bout. The chapter will highlight the network of intracellular pathways (including mTOR signaling) that ultimately lead to increases in gene expression and protein synthesis. Accumulation of acute exercise responses by systematic training over time modulate muscle proteome that can be observed as changes in skeletal muscle phenotype.

**Key words:** Resistance exercise, muscle growth, neuromuscular performance, signaling pathways

(Body text)

Resistance exercise training (RT) can be functionally defined as the progressive overload of a skeletal muscle that is characterized by high muscle contraction force and anaerobic ATP resynthesis. Long-term RT elicits a range of physiological adaptations that contribute to changes in muscle function. Specifically, RT stimulates adaptive machinery responsible for increased maximal contractile force output that is primarily promoted by the combined effect of enhanced muscle activation and muscle fiber hypertrophy<sup>1</sup>. Skeletal muscles play an essential role in locomotion and in the control of whole body metabolism and, hence, RT is widely employed by athletes to improve sport-specific performance, by general population to promote health, lean body mass and general fitness, and in rehabilitation to prevent loss of muscle mass and strength in pathological states<sup>2,3</sup>.

Skeletal muscle can exhibit remarkable plasticity in response to RT. Adjustments within the muscle milieu to mechanical and metabolic demands of RT act to attenuate cellular stress during subsequent exercise sessions<sup>4</sup>. The major morphological adaptation that is specific to RT is a marked increase in muscle cross-sectional area (CSA) of exercised muscles. However, the design of the RT program (i.e. volume, intensity, and frequency of RT sessions, mode of muscle actions used, progression, periodization, variety of RT stimulus, and integration of RT to other fitness training) specifically influences the subsequent chronic physiological adaptations<sup>5</sup>. Also factors such as heredity, sex, biological age, exercise training history, health status and possible medication, diet and nutritional supplements, personal lifestyle and habitual physical activity/inactivity, other physical and psychological stressors (e.g. work-related) and coping with mental stress, training adherence as well as environmental factors can influence chronic outcomes of RT<sup>4,6</sup>.

Changes in muscle size

Increases in skeletal muscle size (i.e. hypertrophy) are the most striking adaptations that occur in response to RT. Muscle hypertrophy is detectable after few weeks from the beginning of the regular RT and proceeds in a linear manner at least for the first few months of training<sup>1</sup>. Typically, on average 6-9 % increases in muscle size are observed in quadriceps muscles following few months of RT in previously untrained individuals.<sup>7</sup> Reaching the individual maximum in muscle mass gains, however, may require several years of systematic RT.

In research, greater hypertrophy is typically observed in upper body muscles compared to lower extremity muscles, possible due to higher locomotive activity in leg muscles that may reduce the

potential for further muscular responses induced by the exercise stimulus<sup>1</sup>. Age and sex may not have substantial effects on the initial RT-induced muscle hypertrophy<sup>1, 8, 9</sup>. However, in women and in older individuals muscle hypertrophy may remain only modest over the years of systematic RT when their responses are compared to those observed in younger adult men. RT-induced muscle hypertrophy occurs specifically in trained muscles and is depending on the design of the RT program. In an untrained state, various kinds of training regimens can induce muscle hypertrophy during the first months of training<sup>10</sup>. For maximizing muscle growth by long-term RT, however, the RT program should principally include multiple sets per exercise with 6-12 heavy repetitions per set and relatively short rest intervals between the sets. Furthermore, several different kinds of exercises should be employed per muscle group in the context of split training routine<sup>5, 11, 12</sup>.

#### Changes in myofiber size and fiber type transitions

Skeletal muscle hypertrophy induced by RT occurs through an increased protein content of individual muscle fibres. Most of the skeletal muscle fibre cytoplasm is occupied by myofibrils with the most abundant proteins being myosin and actin filaments. Thus, RT-induced fibre hypertrophy is primarily a result of the biosynthesis and accumulation of new contractile myofilaments, with concomitant expansion of fibre volume<sup>13-15</sup>. The possible mechanism of fiber growth is an increased CSA and proliferation of myofibrils that results in increased contractile material arranged in parallel and, consequently, an increase in force production capacity. These increases appear especially in fast-twitch type II fibres following RT<sup>1</sup>.

Skeletal myofibers develop distinct phenotypic characteristics during the postnatal period, and therefore the distribution of fiber types is genetically determined. Muscle fibers are characterized as type 1 (slow-twitch) and 2A and 2X (fast-twitch) according to the predomination of their myosin heavy chain (MHC) polymorphisms. Although the fiber-type profile is genetically established, fibers may be remodeled throughout the lifespan by diverse physiological stimuli such as the training status<sup>16</sup>. Many human studies have demonstrated that prolonged RT promotes the conversion of muscle fibers from the glycolytic type 2X fibers to the more oxidative type 2A fibers, whereas the proportion of type I fibres remains rather unchanged<sup>17</sup>. During fiber-type transformation, the expression of MHC isoforms and mitochondrial density, oxidative phosphorylation (OXPHOS) activity, vasculature, and fatigue resistance are switched accordingly. The ability to "shift" skeletal muscle fibers from a more glycolytic fiber to a more oxidative fiber is critical for energy availability to support skeletal muscle contractile activity during the exercise<sup>18</sup>.

#### Changes in muscle architecture

Muscle architecture is defined as the structural arrangement of muscle fibers and connective tissue elements within the muscle relative to the line of force generation at the macroscopic level. Muscle architecture is an important determinant of muscle's mechanical function by affecting the force-velocity relationship.<sup>19</sup> Key components of this relationship are fiber length and pennation angle. Muscle fascicles are bundles of muscle fibers and the angle in which they attach to tendon or aponeurosis (i.e. pennation angle) can be investigated by ultrasound techniques *in vivo*. Physiological CSA (i.e. the area of the cross section of a muscle perpendicular to its fibers) is closely related to the force produced by the muscle and is largely determined by the pennation angle. On the other hand, muscle velocity is proportional to muscle fiber length<sup>20</sup>.

Several studies have shown that RT-induced structural remodeling of the contractile machinery can increase pennation angles to a certain extent in the hypertrophied muscles<sup>21</sup>. These architectural changes in trained muscles affect the manner how force is transmitted from contracting muscle

fibers to tendons and bones. RT-induced increases in pennation angles allows greater contractile material deposition (i.e. addition of sarcomeres in parallel) that further increases physiological CSA and, consequently, enabling an increase in the force generating capacity of the muscle. Mathematically, extensive increases in pennation angle can negatively affect force production of the contracting muscle due to unfavorable arrangement of muscle fibers relative to the axis of force generation to tendon. Thus, when pennation angle increases by excessive hypertrophy, force produced per physiological CSA may decrease<sup>22</sup>. On the other hand, increases in myofibrillar size without changes in muscle architecture (i.e. increase in myofibrillar packing density or lateral force transmission) may explain changes in specific tension (i.e. force exerted by the fibers per unit of physiological CSA). That may partly explain why initial strength gains by RT can be achieved without measurable increase in muscle CSA<sup>22,23</sup>.

#### Satellite cells and myonuclei addition

Skeletal muscle satellite cells are quiescent myogenic precursor cells, located adjacent to muscle beneath the basal lamina but outside the sarcolemma. The role of satellite cells in muscle regeneration is well established. Satellite cells can be activated in response to traumatic lesions requiring muscle regeneration. Once activated, satellite cells proliferate and/or fuse together with preexisting fibers to regenerate muscle tissue<sup>24</sup>.

Also, increased muscle activity during exercise can induce activation and proliferation of satellite cells. Subsequent fusion with an existing myofiber results in the addition of a myonucleus to the fiber syncytium, thereby increasing the total number of myonuclei. The number of myonuclei is a critical determinant of protein synthesis capacity by providing the amount of DNA necessary to sustain gene transcription<sup>25</sup>. The proposed role of the satellite cells in muscle hypertrophy revolves around the concept of a myonuclear domain, meaning that a single myonucleus supports a certain volume of cytoplasm<sup>24,26</sup>. Myofibers are composed of many myonuclear domains and the myonuclear domain size is considered as virtually constant. Satellite cells provide a source for new myonuclei at a rate sufficient to maintain an almost constant myonuclei to-cytoplasmic ratio during skeletal muscle hypertrophy in response to chronic RT. Indeed, several studies have demonstrated that muscle fiber hypertrophy is accompanied by a concomitant increase in satellite cell and/or myonuclear content during chronic RT in humans<sup>25</sup>.

It appears, however, that at least modest muscle hypertrophy can be achieved without the addition of new myonuclei. Thus, the existing myonuclei may have the intrinsic ability to increase their RNA and protein synthesis capacity to such an extent that additional myonuclei are not required to support initial muscle fiber growth<sup>2,27</sup>. However, there may exist a ceiling size of the myonuclear domain area beyond which a fiber will not be able to continue hypertrophy extensively unless more myonuclei are incorporated into the growing fiber<sup>24</sup>. Furthermore, it has been suggested that satellite cells occasionally need to fuse to the muscle fibers to maintain adult muscle mass. It is currently not well known whether other circulating stem cell populations with myogenic potential, together with satellite cells, are activated during the RT-induced hypertrophic response<sup>25</sup>.

Activation of satellite cells and their myonuclear addition has been proposed to play a pivotal role in the phenomenon of "muscle memory" in RT that may be very long lasting in humans. According to the suggested theory, previously untrained fibers recruit myonuclei from activated satellite cells to support hypertrophic growth. During the subsequent detraining and muscle atrophy, myonuclei are protected against the elevated apoptotic activity and the higher number of myonuclei is retained. When muscles are subjected to hypertrophic re-training, muscles grow faster compared to initial RT period. However, convincing evidence of this theory is still lacking in humans<sup>28</sup>.

## Connective tissue adaptations to resistance training

Collagen rich connective tissue constructs tendons and ligaments. In addition, connective tissue exists around and within skeletal muscle. Tendons play a main role in transmitting contractile force to bone and producing elastic energy, while ligaments stabilize joints preventing excessive movements that could damage the joint. Intramuscular connective tissue contributes to passive stiffness in the musculature and force transmission between muscle fibers. Adaptations in the connective tissues occurs specifically only in structures that are exposed to loading<sup>29, 30</sup>.

As skeletal muscles become stronger by RT, intramuscular connective tissues as well as tendons and ligaments also adapt to support increased muscular strength by increasing CSA<sup>31, 32</sup>. The turnover and, thus, renewal of tissue is somewhat slower in connective tissue in the musculoskeletal system compared to that of contractile proteins in the skeletal muscle cells itself. Therefore, tendons appear to hypertrophy at a somewhat slower rate compared to muscles by RT<sup>33, 34</sup>. However, increases in tendon stiffness can occur before tendon hypertrophy. That is possibly due to adaptations in internal structures of the tendon, e.g. modulation of cross-link composition between collagen molecules that improve mechanical properties of the tendon<sup>35, 36</sup>. Increased tendon stiffness may enhance the utilization of elastic energy during stretch shortening cycles and increase the rate of force development during the explosive (fast) muscle actions<sup>33, 37</sup>.

Bone mass, density and architecture are modified to sustain strains produced by the mechanical load. Osteocytes are involved in the turnover of bony matrix through various mechanosensory mechanisms. Bone adapts to external stress specifically at the loaded sites when the magnitude of peak strain is adequate. The most effective intervention for improving bone mineral density (BMD) appears to be exercise characterized by relatively large loading magnitudes and rates. Long-term RT of sufficient intensity and volume has been shown to increase BMD<sup>38-41</sup>.

## Metabolic adaptations to resistance training

Heavy resistance exercise (i.e. several sets with submaximal loads and short rest periods between the sets) can induce acute decrease in ATP, phosphocreatine (PCr) and glycogen storages and marked increase in the concentration of blood lactate, indicating a high rate of anaerobic glycolysis. Consequently, depleted glycogen storages may partly induce post-exercise muscle fatigue<sup>42-44</sup>.

Depending of the exercise-induced myofibrillar disruptions and volume of the loading protocol used and the content of carbohydrates in diet, glycogen storages may be replenished within a couple of days following exercise<sup>45-48</sup>. As a result of chronic RT, the activity of anaerobic enzymes (e.g. creatine phosphokinase, myokinase, and phosphofructokinase) has been shown to increase. Furthermore, intramuscular PCr and glycogen concentrations increase<sup>49, 50</sup>. Chronic RT may also increase skeletal muscle oxidative capacity<sup>51, 52</sup>. These adaptations may lead to improvements in energy metabolism and especially in glycolytic capacity in trained muscle by long-term RT<sup>2</sup>.

Skeletal muscle is the main tissue for glucose disposal accounting for up to 80% of insulin mediated glucose uptake in the postprandial state. During the resistance exercise bout, glucose uptake increases significantly in loaded muscles<sup>51, 53</sup>. Research evidence suggests that chronic RT is effective in improving insulin sensitivity when the intensity is above 50% of 1RM and adaptations occur predominantly locally in the trained skeletal muscles<sup>54, 55</sup>. One of the main mechanisms behind RT-induced muscular adaptations of improving insulin sensitivity is thought to be an increased glucose transport into the myocytes by increased glucose transporter type four (GLUT 4) production, which is the primary transporter facilitating diffusion of circulating glucose into the skeletal muscle cells<sup>56,</sup>

Capillary growth likely occurs in parallel with enlargement of muscle fibers and, thus, capillary density is predominantly maintained in hypertrophied muscles with RT<sup>58</sup>. Mitochondrial content is mainly maintained or reduced following RT<sup>59</sup>. Chronic RT may induce no change or improvements in blood pressure<sup>60-62</sup> and blood lipid profile, of which reductions in LDL cholesterol appears to be a recurrent finding with RT<sup>63-65</sup>. In terms of cardiac morphological changes, resistance-trained athletes may have normal internal diameters but significantly thicker left ventricular wall, referred as to a concentric hypertrophy, although the research findings are not consistent<sup>66-69</sup>. Generally, long-term RT is beneficial for body composition by inducing increases in fat-free mass and decreases in fat mass, which may result in an increased resting metabolic rate<sup>70-72</sup>.

### Molecular adaptations to resistance training

Skeletal muscle demonstrates a remarkable malleability to respond and adapt to contractile activity. The physiological stress by a resistance exercise bout is thought to disrupt cellular homeostasis. Cells react to stress by altering cellular functions to restore homeostasis during and after the exercise bout. Repeated disruptions of homeostasis, followed by sufficient recovery, generate gradually structural and functional adaptations in muscle tissue associated with long-term RT ("progressive overload principle")<sup>73,74</sup>.

From a molecular perspective, RT adaptations are based on the accumulation of specific proteins that alter cellular properties. Remodeling of skeletal muscle starts by loading-specific stimuli that affect the activation of a complex network of intracellular signaling pathways. These signals mediate alterations in enzyme activities, gene expression, and protein biosynthesis, which finally will modulate muscle proteome (Figure 4.1). Thus, the functional outcomes of RT, such as muscle mass gains and metabolic improvements, are coupled to the specificity of molecular responses<sup>2,75</sup>. Although acknowledging that the molecular network regulating skeletal muscle adaptations to RT is vast and affected by numerous factors, this chapter only briefly reviews a few mechanisms that have been suggested to be key players in adaptation to RT in humans.

### Stimuli for muscle adaptations to resistance training

#### Mechanical stress

The mechanical tensile stress (stretch) is one important stressor associated with RT. All forms of muscular activity, whether eccentric or concentric, result in tension (force) through the active muscle(s). Tension associated with RT disturbs the integrity of skeletal muscle, causing mechanochemically transduced molecular and cellular responses (mechanotransduction), favoring anabolism in myofibers and satellite cells. Mechanical stress can induce intracellular responses also independently of growth factors<sup>76-79</sup>.

Mechanical stress induced by high force contractions during the resistance exercise is sensed in loaded muscles by various mechanisms. Mechanical stress induces the release of growth factors, such as IGF-I and hepatocyte growth factor (HGF), from the extracellular matrix. HGF can interact with satellite cells and activate signaling events leading to satellite cell proliferation<sup>80-82</sup>. Muscle contractions transiently disrupt the sarcolemma (the lipid bilayer that surrounds a muscle cell) integrity, which increases the concentration of membrane lipid phosphatidic acid (PA), leading to the activation of signaling pathways inducing hypertrophy<sup>83-85</sup>. Mechanical stress generated in sarcomeres is transferred to the extracellular matrix via costameres, which is a protein complex that connects peripheral myofibrils via the z-disks with the sarcolemma. The costamere comprises a dystrophin/glycoprotein complex and focal adhesion complex, which includes the transmembrane receptor integrin. Activation of integrin can induce hypertrophic signal transduction pathways

through focal adhesion kinases (FAK) <sup>86-90</sup>. The acute increase in intracellular hydration (cell swelling) may occur during the resistance exercise due to lactate accumulation, contributing to the osmotic gradient in skeletal muscle. Increased intracellular pressure may threaten the structural integrity of the cell membrane and, thus, initiate anabolic intracellular signaling response via activation of integrin and focal adhesion proteins <sup>91-93</sup>. Titin is a flexible intrasarcomeric protein that contributes to force transmission and defines the passive stiffness of skeletal muscle. Titin is a likely candidate to sense alterations of mechanical load and interact with diverse cellular signaling pathways inducing hypertrophy <sup>94, 95</sup>. Stretch activated channels (SACs) are calcium and sodium permeable channels which open due to mechanical stress to the sarcolemma. SACs may act as mechanosensor by allowing an influx of calcium ion ( $\text{Ca}^{2+}$ ) into the myofiber that activates  $\text{Ca}^{2+}$  dependent hypertrophic intracellular signaling pathways <sup>96-98</sup>.

### Metabolic stress

In addition to mechanical stress, also metabolic stress associated with RT can promote muscle adaptations <sup>99</sup>. A high rate of ATP turnover during muscle contractions and consequent accumulation of AMP, the release of  $\text{Ca}^{2+}$  from the sarcoplasmic reticulum, and local hypoxia in exercising muscles may stimulate energy-sensing signaling pathways regulating energy metabolism during the resistance exercise <sup>2, 74, 76</sup>. Generally, metabolic stress increases during resistance exercise in an intensity/volume-dependent manner. Presumably, high metabolic stress together with high mechanical strain could be achieved by a hypertrophy-oriented resistance exercise protocol of 6-12 repetitions per set with each set performed until failure, and with relatively short rest periods between the sets <sup>100, 101</sup>.

Hypoxia. Local tissue hypoxia may occur during resistance exercise. Hypoxia is a major cellular stressor and oxygen sensing is well established in the regulation of adaptive processes in cells <sup>102</sup>. HIF-1 is considered as the master regulator of signal transduction pathways sensitive to the changes in intracellular partial pressure of oxygen ( $\text{PiO}_2$ ). Activation of HIF-1 induces transcription of target genes involved in promotion of glucose metabolism and glycolysis, erythropoiesis, and angiogenesis <sup>2, 74</sup>.

REDOX state. Skeletal muscle significantly generates reactive oxygen species (ROS), reactive nitrogen species (RNS), and nitric oxide (NO) during contractile activity. The generation of ROS is dependent on cellular antioxidant capacity and ROS regulates its own defense by promoting cellular antioxidants to maintain redox balance at rest and during the exercise <sup>103, 104</sup>. During resistance exercise, ROS production is likely increased, and it has been suggested to be an important signal in muscle remodeling to a more oxidative phenotype <sup>105, 106</sup>. In the context of human muscle hypertrophic responses to RT, NO may mediate the activation of satellite cells and ROS, if not expressed excessively, mediate activation of several intracellular signaling pathways, such as IGF-I and Mitogen-Activated Protein Kinase (MAPK) cascades, that could be required for muscle growth <sup>82, 107, 108</sup>.

Calcium flux. Calcium facilitates the cross-bridge interaction between myosin and actin filaments during myofibrillar contraction. During muscle contractions, amplitude and duration of calcium oscillations depends on the level of force output by the muscle. Alterations of intracellular  $\text{Ca}^{2+}$  concentrations in myofibers modulate signaling activity of calcineurin and calmodulin-dependent protein kinase (CaMKII).  $\text{Ca}^{2+}$  signaling influences glucose transport, lipid uptake and oxidation, and regulates activity dependent muscle gene expressions that alter fiber type distribution by promoting slow fibre formation <sup>109, 110</sup>.

## Myofibrillar disruptions and inflammation processes

Resistance exercise results at least to some extent in localized muscular damage, of the loaded muscles, depending on the volume and intensity of the exercise and the training history of the participant. Regular RT typically causes mild muscle damage (myofibrillar disruptions) and full recovery normally occurs within a few days<sup>111</sup>. Misalignment of the myofibrils and Z-line streaming are common characteristic following resistance exercise, especially if the training regimen involves predominantly eccentric muscle actions. It has been proposed that due to the specific neural activation strategy for eccentric contractions, as compared to concentric, fewer motor units are recruited for a given load. This would result in a greater requirement of force per active motor unit, predisposing recruited muscle fibres to disrupt especially following unaccustomed eccentric exercise<sup>112-114</sup>. Eccentric training, however, has been shown to be especially effective in promoting muscle hypertrophy and strength gains<sup>115</sup>.

Resistance exercise-induced myofibrillar disruptions are likely caused by physical stress upon the muscle fibres (i.e. mechanical stress) and the result of metabolic deficiencies, possibly through the loss of  $\text{Ca}^{2+}$  homeostasis (i.e. metabolic stress)<sup>111</sup>. Exercise-induced myofibrillar disruptions may elicit a local acute inflammatory response to promote clearance of damaged tissue and muscle regeneration. Inflammatory responses result in movement of fluid, plasma proteins and leukocytes to the site of cellular disruption that is manifested as delayed onset of muscle soreness, muscle stiffness and swelling, and transient decrease of force-generating capacity in loaded muscles<sup>116-118</sup>.

The infiltration of immune cells to the damaged muscles may be observed within 2–3 days following injury. The inflammatory response promotes clearance of damaged tissue and the regeneration of the damaged muscle fibres. The acute inflammatory response after the exercise includes infiltration of neutrophils and local release of proinflammatory cytokines, such as IL-1, IL-6 and TNF-alpha, which mediate protein breakdown<sup>119</sup>. Released cytokines may also stimulate proliferation of satellite cells<sup>120</sup>. Subsequent secondary inflammation includes a significant infiltration of monocytes (i.e. precursors to macrophages) within the damaged muscle fiber to induce further phagocytic activity. Eventually, muscle tissue remodeling occurs during the regenerative phase following the exercise<sup>111, 121</sup>.

## Systemic and local mediators of muscle plasticity: Hormones, growth factors and myokines

Systemic hormones such as testosterone, growth hormone (GH), insulin, insulin-like growth factor (IGF-1) and cortisol can affect muscle mass and growth throughout lifespan. Depending on acute program variables, resistance exercise can elicit transient post-exercise increase in circulating hormone concentrations<sup>122, 123</sup>. Endocrine responses during the exercise may be primarily due to regulation of whole body energy metabolism providing glucose and free fatty acids for energy in working muscles<sup>124, 125</sup>. Hormones can induce their cellular effects through receptor interactions and systemic elevations of circulating hormones presumably increase the likelihood of interactions with receptors located within the muscle tissue and, thus, potentially effects on the responses of RT<sup>123</sup>. However, normal physiological fluctuations in hormones appear to play a preserving, rather than stimulatory, role in the regulation of muscle size<sup>126</sup>. In many studies, resting hormonal concentrations have not shown a significant change during RT despite increases in muscle strength and hypertrophy<sup>122, 123</sup>. However, changes in testosterone and cortisol concentrations may reflect in some cases changes in training load and, thus, overall stress and recovery status in athletes<sup>127</sup>.

The anabolic effects of IGF-I in skeletal muscle have been clearly demonstrated but systemic IGF-I has only a limited influence on the hypertrophic response. However, local (autocrine/paracrine)



expressions of growth factors, such as IGF-I, within the loaded muscles in response to resistance exercise play probably an important role in skeletal muscle hypertrophy. IGF-I induces proliferation of satellite cells and enhance contractile protein accumulation in myofibers by activating IGF-I receptors in the cell membrane<sup>24</sup>. In contrast to IGF-I, myostatin is a major negative regulator of skeletal muscle growth. Myostatin is classified as a “myokine”, i.e. it is produced and secreted by contracting muscle fibers, and subsequently exert auto-, para- and/or endocrine effects. Myostatin inhibits satellite cell activation, repress expression of myogenic regulatory factors and promote proteolysis. Myostatin effects target cells through the Smad2/3 signaling cascade by binding to Activin type II receptors<sup>76</sup>. In addition to myostatin and IGF-I, many other extracellular signaling molecules may potentially modulate skeletal muscle phenotype during resistance training<sup>128-132</sup>.

## Amino acids

Protein availability is a potent modulator of acute molecular responses to resistance exercise. Essential amino acids, especially the branched-chain amino acid leucine, can independently stimulate signaling pathways that subsequently increase protein synthesis rates. Thus, essential amino acids not only act as a substrate but also a signal to promote protein synthetic responses<sup>110, 133</sup>. Dietary protein intake appears to stimulate protein synthesis in a dose-dependent and saturable manner. Ingestion of 20 g of an isolated high-quality protein source, or 30 g of protein as part of a mixed meal, has been recommended to achieve maximal protein synthetic response<sup>134, 135</sup>.

Insert Figure 4.1 here

Figure 4.1. Schematic overview of physiological stimuli by resistance exercise leading to adaptive responses. Depending on program variables, resistance exercise results in a specific milieu of mechanical and metabolic stimuli within the contracting muscle as well as systemic and local release of signaling molecules that lead to the activation of networks of signaling pathways and altered activity of cellular enzymes. These resistance exercise stimuli, together with nutrient availability, induce protein synthesis and tissue regeneration following the exercise. By chronic resistance training, positive net protein synthesis leads to muscle hypertrophy.

## Regulation of protein synthesis and degradation

A bout of resistance exercise results in an increased rate of protein synthesis during recovery and a proportionately smaller and briefer increase in the protein degradation rate<sup>136, 137</sup>. An acute bout of resistance training increases skeletal muscle protein turnover for up to 48 h after completion of exercise<sup>138</sup>. Increases in protein synthesis are suggested to be a result of an increased efficiency of translation per molecule of RNA<sup>139</sup>. The positive net protein synthetic response following resistance exercise results in an accretion of muscle protein over time<sup>140, 141</sup>. Protein synthesis must exceed protein breakdown for an extended period (i.e. few weeks) until RT-induced muscle size changes are detectable<sup>142</sup>. It seems that unaccustomed exercise bouts lead to exaggerated local and systemic stress responses (e.g. dysregulated redox balance) and, thus, induce increases in protein synthesis<sup>75</sup>. However, repeated bouts of exercise blunt protein synthetic response, resulting in an attenuated increase in protein synthesis in the trained state<sup>143, 144</sup>.

mTOR signaling. The most well-described mechanism by which dietary protein and exercise modulates skeletal muscle protein synthesis and subsequently fiber hypertrophy, is the mechanistic target of rapamycin complex 1 (mTORC1) signaling pathway<sup>145, 146</sup>. The activation of mTORC1 is mediated via insulin/IGF-1 receptor activation by hormones and growth factors and the activation of the downstream phosphatidylinositol-3 kinase (PI3K) - Akt pathway<sup>147</sup>. mTORC1 can be activated

also independently of Akt via contractile activity (mechanotransduction) and essential amino acid provision<sup>148-150</sup>.

Activation of mTORC1 triggers downstream signaling through p70 ribosomal S6 kinase (p70 S6K1), that is a key regulator of protein synthesis through canonical pathways of protein translation and ribosome biogenesis<sup>151, 152</sup>. p70 S6K1 exerts its effect through ribosomal protein S6 (rpS6), eukaryotic elongation factor 2 kinase (eEF2), and eukaryotic initiation factor 4E-binding protein (4E-BP1), which collectively increases mRNA translational efficiency and ultimately protein synthesis for cellular hypertrophy<sup>153, 154</sup>. Increases in resistance exercise volume has been shown to induce pronounced activation of mTOR signaling proteins<sup>155-158</sup>. (Figure 4.2)

AMPK Signaling. The high rate of ATP turnover during the resistance exercise leads to a cellular energy deficit and increases in the AMP/ATP ratio. AMP-activated protein kinase (AMPK) senses the increase in energy turnover when muscle tissue is activated and acts as a signal transducer for metabolic adaptations<sup>159, 160</sup>. Acute exercise increases AMPK enzymatic activity in an intensity-dependent manner, reflecting effects of exercise on ATP turnover. Furthermore, contraction-induced Ca<sup>2+</sup> release and ROS production leads to the activation of AMPK. As expected, resistance exercise has been shown to acutely increase AMPK activity in skeletal muscle<sup>161-163</sup>.

Overall, AMPK activation acts to conserve ATP by inhibiting biosynthetic and anabolic pathways, while simultaneously stimulating catabolic pathways to restore cellular energy stores. In skeletal muscle during the exercise, AMPK activation modulates cellular metabolism acutely through phosphorylation of metabolic enzymes that suppresses glycogen and protein synthesis, but promotes lipid metabolism and glucose uptake<sup>159, 160, 164</sup>. Protein synthesis is an energy-consuming process and, in agreement with the role as an energy sensor, activation of AMPK can suppress protein synthesis by inhibiting directly mTORC1 activity or indirectly through mTOR upstream kinase tuberin (TSC2) activation<sup>162, 165, 166</sup>. AMPK can also inhibit protein synthesis through activation of eukaryotic translation elongation factor 2 kinase (eEF2K), leading to inhibition of protein translation by eEF2<sup>4, 167</sup>. Like AMPK, also energy sensors REDD1 (regulated in DNA damage and development 1) that is activated by ATP depletion and hypoxia-can inhibit mTORC1 and subsequently protein synthesis<sup>4, 168</sup>. Chronic AMPK activation alters metabolic gene expression and induces mitochondrial biogenesis, leading to promotion of an oxidative muscle phenotype. The long-term regulatory actions are mediated via direct phosphorylation of transcription factors and the transcriptional coactivator PGC-1 $\alpha$ <sup>169, 170</sup>.

Protein degradation by the ubiquitin-proteasome system is regulated via muscle-specific E3 ubiquitin ligases, muscle atrophy F box (atrogin-1/MAFbx) and muscle RING finger 1 (MuRF1). They are involved in ubiquitination of specific proteins that are transferred to the 26S proteasome for subsequent degradation<sup>171</sup>. For example, MuRF1-dependent ubiquitination regulates the degradation of contractile proteins, such as myosin heavy chains<sup>172</sup>. Transcriptional upregulation of atrogin-1/MAFbx and MuRF1 are increased by activation of the forkhead box O (FOXO) family of transcription factors<sup>173</sup>. Anabolic and energy sensitive processes regulate the ubiquitin-proteasome pathway activity. Akt, an upstream mediator of mTORC1 activity, inactivate FOXOs and, thus, inhibits proteolysis. Contrarily, AMPK activation promotes FOXOs activity<sup>110, 174</sup>. Also, inflammatory response via the NF- $\kappa$ B pathway and myostatin promote proteolysis through ubiquitin proteasome system<sup>175-178</sup>.

Autophagy refers to a process of non-selective degradation of cytosolic components by the lysosome. Autophagy is beneficial to maintain cellular homeostasis at rest as well as during the exercise and post-exercise recovery<sup>107, 179</sup>. Resistance exercise may induce certain damage to loaded

myofibers and autophagy enables removal of damaged organelles and proteins through the lysosomes. It appears that adaptation to exercise training is reliant on proper activation of autophagy, and acute physical exercise has been shown to be a potent inducer of autophagy in skeletal muscle<sup>180-183</sup>. Activity of autophagy is regulated through unc-51-like kinase 1 (ULK1). Activated mTORC1 can inhibit autophagy while AMPK stimulates autophagy by regulation of ULK1 activity<sup>162, 179, 184, 185</sup>. Besides the autophagy-lysosomal system, non-lysosomal proteases like the calcium-dependent calpain family and caspase class of proteins are involved in proteolysis<sup>105, 186, 187</sup>. For example, the sarcomeric damage caused by exercise initiates a calpain-mediated degradation of disrupted sarcomeric filaments, such as titin<sup>94</sup>.

Insert Figure 4.2 here

Figure 4.2. Simplified overview of mTOR and AMPK signaling pathways regulating skeletal muscle size and function by resistance exercise. Putative resistance exercise-induced stimuli (amino acids and contractile activity induced mechanic and metabolic stress, and release of systemic and local signaling molecules), activate specific intracellular signaling networks (including, but not limited to mTOR and AMPK signaling) that mediate acute and chronic skeletal muscle responses to resistance exercise training. Stimulation of the signaling pathways depends on the resistance exercise program variables used<sup>155-158, 188-190</sup>. 4E-BP1, eukaryotic initiation factor 4E-binding protein 1; Akt, protein kinase B; AMPK, AMP-activated protein kinase; AS160, Akt substrate of 160 kDa; CaMK, calmodulin-dependent protein kinase; eIF2B, eukaryotic initiation factor 2B; FAK, focal adhesion kinase; FoxO, forkhead box protein; GSK3, glycogen synthase kinase 3; mTORC1, mammalian/mechanistic target of rapamycin complex 1; p38 MAPK, p38 mitogen-activated protein kinase; p70S6K, ribosomal S6 kinase 1; PA, phosphatidic acid; PGC-1 $\alpha$ , peroxisome-proliferator-activated receptor gamma, coactivator 1; Rag, Ras-related small GTPase; RONS, reactive oxygen and nitrogen species; ULK1, Unc-51-like kinase 1. Arrows denote activation, oval arrows denote inhibition.

## Summary

The aim of RT is to provide an overload stimulus across the muscle to generate specific molecular responses, promoting adaptive changes in skeletal muscle mass and metabolic function. The conversion of intra- and extracellular signals generated during muscle contractions to subsequent physiological adaptations involves a cascade of stimuli that affect specific signaling pathways regulating exercise-induced satellite cell activation, gene expression and/or protein turnover rate by proteosynthesis and proteolysis<sup>24, 110, 126</sup>.

RT-induced skeletal muscle hypertrophy appears to be significantly controlled by the activity of the translational machinery, where mTORC1 acts as the master regulator. Several signaling pathways have an integrated effect on mTORC1 activity, of which autocrine/paracrine growth factors, mechanical and metabolic stress and amino acid availability appears to be the most prominent<sup>2, 74, 76</sup>. However, RT-induced skeletal muscle adaptation is a complex process involving various cellular responses and many signal transduction pathways. Cellular signaling pathways comprise complex networks that are operative in constantly altering cellular milieu, making it difficult to connect specific signaling responses and changes in gene and protein expression to certain metabolic responses after a resistance exercise bout, let alone to long-term adaptations to RT<sup>191, 192</sup>. In addition to the transcriptional and translational signaling networks involved with skeletal muscle plasticity, epigenetic modifications of DNA that may affect exercise-induced gene expression, or post-transcriptional silencing of genes by miRNAs, comprise additional levels of control on adaptations to RT<sup>193-197</sup>.

To entirely understand the adaptive changes that myofibers undergo in response to RT in humans, aspects of the neuromuscular system (e.g. neural drive to muscles), musculoskeletal system (e.g. force transmission from muscles to tendons and bones), and cardiovascular system (e.g. muscle capillary network) should be taken into consideration. Furthermore, RT-induced adaptations generally affect the entire body, thus, adaptations in musculature should be viewed in the context of the crosstalk between tissues and organs in a whole body <sup>198-201</sup>.

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