CAN MUSCLE PROTEIN METABOLISM BE SPECIFICALLY TARGETED BY EXERCISE TRAINING IN COPD?

Davina C. M. Simoes\textsuperscript{1*} and Ioannis Vogiatzis\textsuperscript{1}

\textsuperscript{1}Department of Sport, Exercise and Rehabilitation, Faculty of Health and Life Sciences, Northumbria University Newcastle, Newcastle-upon-Tyne, United Kingdom.

*Corresponding author: Davina C M Simoes, Department of Sport, Exercise and Rehabilitation, Faculty of Health and Life Sciences, Northumbria University, Newcastle-upon-Tyne, United Kingdom, email: davina.simoes@northumbria.ac.uk

Running title: Exercise-induced muscle hypertrophy in COPD
Abstract

Patients with stable chronic obstructive pulmonary disease (COPD) frequently exhibit unintentional accentuated peripheral muscle loss and dysfunction. Skeletal muscle mass in these patients is a strong independent predictor of morbidity and mortality. Factors including protein anabolism/catabolism imbalance, hypoxia, physical inactivity, inflammation, and oxidative stress are involved in the initiation and progression of muscle wasting in these patients.

Exercise training remains the most powerful intervention for reversing, in part, muscle wasting in COPD. Independently of the status of systemic or local muscle inflammation, rehabilitative exercise training induces up-regulation of key factors governing skeletal muscle hypertrophy and regeneration. However, COPD patients presenting similar degrees of lung dysfunction do not respond alike to a given rehabilitative exercise stimulus. In addition, a proportion of patients experience limited clinical outcomes, even when exercise training has been adequately performed. Consistently, several reports provide evidence that the muscles of COPD patients present training-induced myogenic activity resistance as exercise training induces a limited number of differentially expressed genes, which are mostly associated with protein degradation.

This review summarises the nature of muscle adaptations induced by exercise training, promoted both by changes in the expression of contractile proteins and their function typically controlled by intracellular signalling and transcriptional responses. Rehabilitative exercise training in COPD patients induces skeletal muscle mechanosensitive signalling pathways for protein accretion and its regulation during muscle contraction. Exercise training also induces synthesis of myogenic proteins by which COPD skeletal muscle promotes hypertrophy leading to fusion of myogenic cells to the myofibre. Understanding of the biological mechanisms that regulate exercise training-induced muscle growth and regeneration is necessary for implementing therapeutic strategies specifically targeting myogenesis and hypertrophy in these patients.

Keywords: COPD, exercise, protein synthesis, anabolism, hypertrophy, myogenesis, muscle wasting
**Introduction**

COPD is the fourth leading cause of death worldwide and is projected to be the third most common cause of death by 2020 (1). Cigarette smoke constitutes the major preventable risk factor, resulting in a progressive proteolytic, inflammatory and vasoactive response that leads to emphysema, small airway obstruction and pulmonary hypertension. The oxidative stress imposed by cigarette smoking together with systemic inflammation and hypoxia are important contributors to pathogenesis of skeletal muscle wasting and dysfunction and have been previously extensively reviewed (2-4). Skeletal muscle wasting is an important systemic effect of the disease and a strong independent predictor of mortality (5-7). This unintentional accentuated skeletal muscle wasting is frequently associated with altered muscle structure (fiber size, fiber type distribution, capillary density and metabolic capacity) and dysfunction (decreased strength and endurance). While the magnitude of the alterations varies substantially across individuals, some degree of muscle wasting affects all individuals during ageing. Age-related defects in protein metabolism have been proposed to be causally involved in this muscle loss (8). These changes are attributed to both inactivity- and age-related alterations in protein synthesis and degradation, indicating complex pathophysiological phenomena involving both structural changes to the muscle fibres, as well as the enzymatic machinery that controls metabolism. Independently of the factors promoting muscle wasting in COPD, regular physical exercise remains the most potent available treatment option for reversing, in part, locomotor muscle wasting and dysfunction in COPD (9). Indisputably, exercise training promotes a range of beneficial adaptations in the skeletal muscle including increased capilarization, fibre type plasticity, hypertrophy and function. However, exercise training induced muscle benefits are certainly much smaller in COPD patients compared to age matched controls. In addition, not all COPD patients respond adequately to exercise stimulus, even when exercise training is properly performed (10, 11). An improved understanding of molecular mechanisms of exercise induced adaptations in COPD and healthy individuals will be valuable to inform future directions to address the issues on resistance to exercise-induced adaptations in COPD.

**Effect of prescribed exercise training to skeletal muscle adaptation in COPD**

Exercise training is a mechanical stimulus that consists of repeated, episodic bouts of muscle contraction promoting functional adaptation and remodelling not only to the skeletal muscle,
but also to various systems in our body (12, 13). Exercise promotes a range of adaptations that is beyond the musculoskeletal system promoting general health (14). Briefly, in parallel with neural signals to the skeletal muscle contraction, powerful neural feed-forward signals to the respiratory, cardiovascular, metabolic and hormonal systems are produced. In response to exercise training, COPD patients demonstrate reduction in dyspnea sensations, improvements in exercise capacity and quality of life (9).

Skeletal muscle adaptations are dependent on the intensity and duration of the exercise training performed (15). High intensity exercise training ($W_{\text{peak}} \geq 80\%$) is generally described to promote improvement in exercise capacity. However, COPD patients with limited ventilatory capacity are usually unable to sustain high intensities for sufficiently long periods. Taking into consideration COPD patients exercise capacity, a number of studies have been performed employing a varied combination of exercise modalities/training, program duration and intensity. Usually, the duration of an exercise training programme is set from 8 to 12 weeks and as frequent as 3 times per week. Combination of high intensity aerobic and resistance exercise is described to promote quantifiable muscle hypertrophy in COPD (16, 10, 17, 11). Conversely, exercise training of lower intensity ($W_{\text{peak}} \leq 60\%$) was found unable to promote quantifiable changes in muscle hypertrophy and fibre type distribution (18, 19).

Therefore, programmes incorporating high intensity exercise training are more likely to induce quantifiable skeletal muscle adaptations.

Different modes of exercise such as endurance- and resistance–based are known to stimulate variable but specific skeletal muscle adaptations, leading to muscle endurance and strength respectively (15). Aerobic/endurance exercise training enhances mitochondrial protein content and oxidative capacity of trained myofibers, improving insulin sensitivity and skeletal muscle metabolic function (20). Whereas, resistance training increases myosin-heavy-chain gene transcripts and synthesis rate of muscle proteins promoting strength (15). When comparing different modalities of exercise prescribed to healthy sedentary people, Robinson et al (21) observed that high intensity interval training (HIIT) enhanced more comprehensively changes such as aerobic capacity, mitochondria respiration and lean body mass (21). HIIT training simultaneously promoted endurance- and resistance–based training skeletal muscle adaptations, that promoted changes in transcription and translation regulation of muscle growth and mitochondrial pathways (21). HIIT reversed age-related proteome, particularly of mitochondrial proteins. But both resistance training and HIIT increase proteins
involved in translational machinery. HIIT exercise involves 30s repeated short bouts of activity at near maximal intensity (Wpeak ≥ 80%) interspersed with 30s rest periods, which despite its high intensity, has been successfully applied to COPD patients (10, 17, 22, 11). A representation of basic skeletal muscle adaptation promoted by aerobic/endurance, resistance and HIIT is depicted in Figure 1, originally adopted by Robinson et al. (21).

Therefore, adaptations seen in COPD skeletal muscle take place in accordance with specific exercise training stimuli, thereby partially explaining the large variability in muscle adaptations seen among studies. The effect of exercise modalities in promoting COPD skeletal muscle adaptations at structural and protein metabolism levels have been recently presented in a systemic review (18).

Histologically, the skeletal muscle appears uniform, but is composed by a range of heterogeneous myofibres regarding size, metabolism and contractile function. When comparing myosin-heavy-chain isoform expression, myofibres are classified into type I, type IIa, type IId/x, and type IIb fibres (23). Type I and IIa fibers exhibit high oxidative potential and IIx and IIb are primarily glycolytic. Whether endurance- or resistance-based exercise training can induce myofibre plasticity is still debatable. Certainly, COPD patients present a shift in fibre type displaying fewer type I (oxidative) fibres and greater proportion of type II (glycolytic) fibres in quadriceps muscles (24). This shifting towards glycolytic fibres is associated with increased mortality (7). Exercise training prescribed to COPD patient can only partially reverse this fibre type shifting. Proportion of fiber type I and IIa were increased mainly after HIIT and high intensive aerobic exercise (18, 25) as endurance training confers an increased oxidative profile to trained myofibers (23). Hypertrophy of fibre type I and IIA was more widespread among different modalities of exercise training, (18), Whereas, capillary to fiber ratio adaptations were though observed across various intensities and modalities of exercise training (18).

**Contribution of inactivity to COPD muscle wasting**

Inactivity appears to be an important mechanism in the process of muscle loss in COPD, given that muscles that are active, such as the diaphragm and the adductor policis, do not exhibit atrophy in contrast to inactive muscles, such as the quadriceps (26). In experiments comparing different muscle groups in COPD patients, the characteristics exhibited by the
deltoid and the diaphragm were different compared to the quadriceps (27). Importantly, the muscles of respiration in COPD exhibit a contrary shift in fiber typing compared to the locomotor muscles, manifested by increased type I fiber distribution (28, 29). Various conditions of reduction in neuromuscular activity promoted by inactivity are known to decrease myonuclear number in atrophying muscle and impact on fibre typing. As reviewed elsewhere (30), detraining experiments in healthy individuals have shown to induce locomotor muscle adaptations that lead to increased number of muscle fibre type IIX phenotype, and attenuation of mitochondrial biogenesis (PPARs and PGC-1α) as observed in COPD patients. As described above, reassuming activity by exercise training can partially promote changes in fibre type distribution (18). However, inactivity alone does not seem to fully explain the phenomenon of muscle fibre type shifting in COPD.

**Mechanical stress and mitogen-activated protein kinase signalling**

The ability of skeletal muscles to respond to physical exercise by executing the appropriate metabolic and transcriptional response is dependent upon the cellular signal transduction through phosphorylation cascades. Multiple kinases, including AMPK, Akt and the mitogen-activated protein kinases (MAPKs) are involved in the regulation of DNA transcription through the phosphorylation of nuclear transcription factors. This either enhances or inhibits the ability of transcription factors to bind DNA, affecting target gene transcription (31, 32). Three main MAPK subfamilies are activated by acute exercise in human skeletal muscle: (1) the extracellular-regulated kinase (ERK1/2), (2) the c-jun N-terminal kinase (JNK), and (3) the p38 MAPK. Activation of MAPKs regulates the transcriptional events by phosphorylation of diverse substrates localised in the cytoplasm or nucleus, including transcription factors, inducing differentiation, hypertrophy, inflammation, and gene expression (33).

The MAPK p38 is a stress-activated kinase that is transiently activated in response to a strenuous range of stimuli such as physical inactivity and increased intensity of exercise training (34, 35). Activation of p38 in skeletal muscle myoblasts is related to loss of cell autonomous self-renewal capacity (36). MAPK p38 activation is also observed during skeletal muscle immobilisation in a rat hind limb model of acute muscle wasting (37, 34). COPD are generally more inactive compared to their age-matched healthy counterparts (38).
Accordingly, ratios of phosphorylated to total level of p38 MAPK and ERK 1/2 were significantly elevated in patients with COPD compared to controls (39). Whereas, another study have shown no differences in the ratio of phospho-p38 MAPK to total level of p38 MAPK protein between COPD patients and healthy age-matched donors. Although patients with COPD present muscle wasting, discrepancies among studies would be expected as it is uncertain whether patients are actively losing muscle mass at the time of experimentation.

**Major signaling pathways involved in the control of exercise training induced skeletal muscle adaptations**

Endurance- and resistance-based modalities of exercise are controlled by two major signaling pathways regulating mitochondria biogenesis and hypertrophy respectively, as depicted in Figure 2.

The regulation of mitochondrial biogenesis by endurance-based exercise converge from activation of the cascades AMPK and p38 upregulating PGC-1α. When compared to healthy controls, mitochondria density is lower in quadriceps muscle of COPD patients, presenting lower expression of peroxisome proliferator-activated receptors (PPARs), PPAR-γ co-activator 1α (PGC-1α) and mitochondrial transcription factor (TFAM) in cachectic COPD (40). In patients with COPD, exercise enhances the decrease in mitochondria DNA content of skeletal muscle and the expression of PGC-1α mRNA seen in healthy subjects, probably due to oxidative stress (41).

In contrast, resistance training is described to stimulate the signaling pathways responsible for muscle hypertrophy (12, 13). The activation of mTOR and IGF-I appears to be important in this process. To restore muscle mass via regular exercise training, protein synthesis should exceed protein breakdown over an extended period. Hypertrophy of skeletal muscle as result of resistance exercise training is strongly associated with the degree of mTOR activation, ribosomal protein S6K (p70S6K) phosphorylation and downstream targets (42). Contraction-induced p70S6K activation is dependent on mTOR activation, which increases protein translation and inhibits protein degradation via inhibition of both ubiquitin proteasome (43, 32, 44) and autophagy-lysosome pathways (45, 46). mTOR activation is critical to load-
induced muscle growth, as demonstrated by the attenuation of hypertrophy responses and protein synthesis by the mTOR inhibitor, rapamycin (43).

We and others have described that Akt/mTOR pathway is downregulated in skeletal muscle of patients with COPD compared to healthy subjects (10). HIIT promotes the activation of the Akt/mTOR pathway in skeletal muscle only in COPD patients with preserved muscle mass compared to aged matched controls. Some studies suggest that hypoxemia characteristically observed in more severe cachectic COPD patients, is associated with resistance of skeletal muscle activation of the Akt/mTOR pathway (16). Exercise training in hypoxemic patients with COPD was not capable to promote muscle fibre hypertrophy and activation of the Akt/mTOR pathway as compared to normoxemic COPD patients (16). Both in vitro C2C12 myotubes cultured in normoxic and hypoxic conditions and mice models of hypoxia suggest that the response of the Akt/mTOR pathway to exercise could be compromised in hypoxemic patients (16, 47). Therefore, impairment of skeletal muscle hypertrophy commonly linked to the severity of the disease is associated with the magnitude of muscle wasting, the degree of hypoxia, or both. Interestingly, induced expression of the adaptive response of hypoxia HIF-1 responsive RTP801 (DDIT4) is observed only in trained COPD and not in healthy subjects (48).

mTOR regulates the mechanisms of protein synthesis at several levels (e.g. translation capacity, translation efficiency) through increases of translation of specific mRNAs, which culminates in skeletal muscle fibre enlargement. mTOR exists as part of two multi-protein complexes: i) mTORC1, which contains raptor and confers rapamycin sensitivity, is required for signalling to p70S6K and 4E-BP1, whereas ii) mTORC2, which contains rictor and is rapamycin insensitive, is required for signalling to Akt-FOXO (49). The effect of mTOR activity on downstream regulators of protein synthesis is principally achieved through a contraction-induced regulation of mTORC1 (50).

Early work on adaptive hypertrophy has focused on the (transient) post exercise rise in blood-borne anabolic hormones, such as growth hormone and insulin-like growth factor-I (IGF-I), and the consequent activation of the muscle protein synthesis of a signalling cascade (phosphatidylinositol 3-kinase (PI3K)-Akt-mTOR) by IGF-I interaction with insulin and IGF receptors (51). Recently, the muscle growth paradigm has shifted focus to IGF-I-independent mechanisms of mTOR activation and adaptive hypertrophy through mechano-
sensory regulation (50). Nutrient-dependent regulation of muscle growth is achieved through insulin- and Akt-dependent activation of the mTOR pathways. These pathways operate synergistically causing muscle growth and can be augmented by appropriate nutritional intake such as post exercise carbohydrate and amino acid ingestion or increased dietary protein (52). Therefore, IGF-I is involved not only in hypertrophy but also in promoting myogenesis and muscle regeneration.

**Regulation of skeletal muscle protein synthesis promoting myogenesis and muscle regeneration**

Skeletal muscle hypertrophy is achieved by both positive protein balance and fusion of satellite cells to myofibres (Figure 1). It is a process that involves (1) accretion of protein in various cellular compartments via mechanosensitive signalling pathways that drive translation, and (2) the activation and recruitment of resident muscle stem cell (satellite cells) that differentiate to fusion-competent myoblasts (53, 54). Regulation of protein translation and synthesis promotes accretion, whereas activation and incorporation of satellite cells facilitates the addition of myofibrils to the muscle.

Myogenesis and muscle regeneration depend on critical steps for activation of quiescent satellite cell, proliferation, migration, differentiation, fusion and maturation. Fusion of a satellite cell to an existent myofibre results in an increase in the number of myonuclei, and thus the available total amount of genetic machinery for protein production (54). The average number of myonuclei per muscle fibre of non-cachectic COPD is twice as high compared to controls, indicating higher capacity of protein metabolism necessary for maintenance of muscle mass. Quiescent satellite cells are essential to replenishment of myonuclei pool. As satellite cells replicate throughout lifespan, telomeres are shortened. Telomere shortening is a marker of senescence. COPD patients, despite the observation that satellite cells numbers are unaltered in the limb muscle compared to controls, satellite cells present shorter telomeres. A fact suggesting exhausted muscle regenerative capacity, compromising the maintenance of muscle mass (55).

Satellite cells myogenic regulatory factors (MRF) and myostatin play important roles in myogenesis and muscle regeneration. mRNA and protein expression of the myogenic differentiation factor D (MyoD), involved in the proliferation process, has been shown to
increase in the skeletal muscle of non-cachectic COPD patients after HIIT, but no changes were observed in cachectic COPD patients after HIIT (10). MyoD protein expression was also increased after resistance training with or without nutritional supplementation in patients and healthy controls (56). mRNA expression of MRF myogenin, involved in the differentiation process, was not different between a resistance training group and a control group, while addition of testosterone supplementation to resistance training increased myogenin mRNA expression compared to the control group (57). mRNA and protein expression of the myogenic inhibiting factor myostatin, showed no significant change after resistance exercise training neither in patients nor in healthy controls, or after combined aerobic and resistance training (56, 16). After HIIT however, non-cachectic COPD patients showed a significant decrease in mRNA and protein expression of myostatin, while in cachectic COPD patients HIIT did not significantly alter mRNA or protein myostatin expression (10). In addition, mRNA expression of a negative regulator of cell proliferation Kruppel-like factor 10 (KLF11) was increased after aerobic training in patients, but not in healthy controls (48).

The IGF-I system plays an important role in the regulation of muscle cell growth, muscle cell proliferation and muscle cell survival (58, 59). After HIIT, mRNA expression of both IGF-I and the mechano-growth factor (MGF), an isoform of IGF-I, significantly increases. Both cachectic and non-cachectic COPD patients have shown enhanced MGF mRNA expression post-training (10). However, no significant increase in protein expression of IGF-I and MGF expression was observed in cachectic COPD patients in comparison to non-cachectic. Other exercise protocols such as combined aerobic and resistance training were not capable to increase the expression of the IGF-I variants (16). However, testosterone supplementation was found to increase the levels of IGF-I and MGF protein expression in COPD patients after resistance training (57).

Exercise induced changes in gene expression

Comparing gene transcription among different modalities of exercise in healthy individuals, Robinson et al (21) found that HIIT promotes a stronger increase in gene transcripts than other modalities of exercise tested, particularly in older adults.
Skeletal muscle gene expression from COPD after high intensity aerobic exercise training compared to age-matched controls were analysed using Genechip Array (48). High aerobic exercise training induced up-regulation of quantitatively significantly fewer genes in the skeletal muscle of COPD (107 were upregulated and 124 were downregulated) compared with healthy controls (258 were upregulated and 315 were downregulated). Qualitatively, genes associated with protein degradation, such as oxidative stress, ubiquitin proteasome, and COX pathways were distinctly induced only in patients with COPD, potentially reflecting the specific molecular response of the muscle to exercise in COPD, thereby suggesting additional operating mechanisms for exercise limitation in these patients. Whether exercise training can sufficiently enhance muscle hypertrophy to outstrip muscle wasting in COPD patients with substantial muscle loss, remains an unresolved issue.

**Regulation of skeletal muscle protein breakdown**

Muscle tissue homeostasis is maintained by a tight and complex balance between protein synthesis and degradation. Protein metabolism turnover is a dynamic process balancing protein synthesis and breakdown. Muscle wasting due to an increase in protein breakdown is a feature shared among many acute and chronic disease entities as well as healthy ageing.

Muscle wasting has primarily been attributed to increased protein degradation. Protein degradation in COPD patients peripheral muscles takes place through four proteolytic systems, including the ubiquitin-proteasome (UP) pathway, the calpain pathway, the caspase pathway, and the autophagy-lysosome (AL) pathway as reviewed elsewhere (60).

Exercise has been found to stimulate mitogen activated protein kinases MAPK-9 and MAPK activated protein kinase 3 (MAPKAPK-3) in COPD compared to healthy controls (48). The MAPK pathway, in turn activates forkhead transcription factors, involved in muscle protein degradation. When inflammatory response to exercise is limited, the muscle recovers in a timely manner; however, persistent systemic inflammation described in COPD, may be associated with muscle wasting and adversely impact on muscle protein metabolism (61). Numerous pathological indicators in COPD, namely systemic inflammation, hypoxia and oxidative stress most likely trigger catabolic processes in skeletal muscle, that are mediated by transcriptional regulators including nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kB) and forkhead box O transcription factors (FOXOs). The activity of NF-kB is
increased in COPD compared with healthy age-matched individuals (18, 62, 10, 22) and in particular in patients with muscle wasting compared to those without muscle wasting (63, 10). FOXO mRNA and protein expression is increased in patients with COPD (64-67, 62). The expression of FOXO-1 may be associated with physical inactivity as protein expression is increased in lower limbs compared to respiratory muscles in COPD patients, but not in healthy controls (68). Increased catabolic signaling through FOXO and NF-κB activation may induce gene expression of key factors in both ubiquitin proteasome system (UPS) (69, 32) and the autophagy lysosome pathways (45).

**Concluding remarks**

Exercise training promotes a range of beneficial adaptations in the skeletal muscle including increased capilarization, fibre type plasticity, hypertrophy and function. All these adaptations are a result of exercise stimuli that challenges muscle homeostasis by activating networks of signalling molecules. Activation of kinases and pre-transcriptional regulation occurs rapidly during exercise and recovery, whereas protein transcription is subsequently regulated. Intensity, duration, and mode of the exercise stimuli collectively contribute to the relative activation and the magnitude of activated pathways and downstream targets (12, 13). All these parameters have to be considered when designing exercise studies, so results are comparable and can advance knowledge in the area. Future studies on the molecular mechanisms of exercise induced satellite cell myogenic capacity in COPD patients are fundamental for designing pharmacological and exercise training interventions aiming to address resistance to exercise-induced adaptations.

**Acknowledgment:** DCMS is supported by University of Northumbria Multidisciplinary Research Theme (MDRT).

**Author contribution:** DCMS and IV wrote the paper, DCMS and IV approved the final version of the manuscript.


Exercise training is a powerful stimulus producing hypertrophy and regeneration of muscle by increased protein metabolism and fusion of satellite cells to existent myofiber. Endurance- and resistance-based exercise training programmes are characterised as stimuli capable for increasing oxidative capacity and hypertrophy, respectively. Skeletal muscle adaptations observed from combined endurance/resistance as well as high intensity interval training reflect a wider range of stimuli. (The Figure was originally adopted by Robinson et al. (21) and subsequently modified by authors).
Figure 2: Diagram of the major signalling pathways involved in the control of skeletal muscle hypertrophy and mitochondrial biogenesis. Voluntary exercise training activate kinases/phosphatases to mediate a specific exercise-induced signal. The cross talk among the numerous signalling pathways activated and the multiple site regulation produces a high sensitive and complex transduction network.