

## COMPENDIUM ON INTERORGAN CROSSTALK IN HEART FAILURE AND CARDIOMETABOLIC DISEASES

# Skeletal Muscle as a Mediator of Interorgan Crosstalk During Exercise: Implications for Aging and Obesity

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**ABSTRACT:** Physical exercise is critical for preventing and managing chronic conditions, such as cardiovascular disease, type 2 diabetes, hypertension, and sarcopenia. Regular physical activity significantly reduces cardiovascular and all-cause mortality. Exercise also enhances metabolic health by promoting muscle growth, mitochondrial biogenesis, and improved nutrient storage while preventing age-related muscle dysfunction. Key metabolic benefits include increased glucose uptake, enhanced fat oxidation, and the release of exercise-induced molecules called myokines, which mediate interorgan communication and improve overall metabolic function. These myokines and other exercise-induced signaling molecules hold promise as therapeutic targets for aging and obesity-related conditions.

**Key Words:** cardiovascular diseases ■ epinephrine ■ hypertension ■ muscle, skeletal ■ sarcopenia

Physical exercise is widely recognized as an effective tool for preventing, managing, and treating a variety of chronic conditions, including cardiovascular disease (CVD), type 2 diabetes, hypertension, obesity, and sarcopenia.<sup>1,2,3</sup> A systematic review and meta-analysis of 33 studies involving over 880 000 participants revealed that higher levels of physical activity were linked to a 30% to 50% reduction in cardiovascular mortality and a 20% to 50% reduction in all-cause mortality.<sup>4</sup> In addition, research from the Nurses' Health Study (with nearly 80 000 participants) and the Health Professionals Follow-Up Study (involving 44 000 participants) examined the impact of 5 lifestyle factors, including at least 30 minutes of moderate to vigorous physical activity per day, on life expectancy in the US population. Over a follow-up period of up to 34 years, the most physically active men and women enjoyed an increase in life expectancy of 7 to 8 years.<sup>5</sup>

Exercise enhances metabolic health by inducing adaptations across multiple tissues, including skeletal muscle. In skeletal muscle, regular resistance exercise

increases myocyte size<sup>6</sup> and muscle mass<sup>7</sup> and improves the quality and functionality of the muscle.<sup>8</sup> This is due to changes in fiber type and increased mitochondrial content, conferring resistance to atrophy,<sup>9</sup> as well as metabolic adaptations. Endurance exercise results in extensive mitochondrial biogenesis, enhanced glucose and fatty acid transport, and increased capillarization to improve the flux of oxygen delivery and nutrient flux to the skeletal muscle. Nutrient availability is also improved through greater intramuscular lipid and glycogen storage.<sup>10,11</sup> Exercise protects the skeletal muscle from age-associated dysfunction, including decreases in strength and mitochondrial capacity, and increased fat infiltration and insulin resistance.<sup>11,12,13</sup> Moreover, maintaining or beginning physical activity is a well-established way to prevent or improve the effects of aging on skeletal muscle.<sup>14</sup> Exercise results in multiple signals that change skeletal muscle function and metabolism, catecholamine and adrenaline signaling, calcium release, mechanical force, and changes in redox balance that all work to adapt the skeletal muscle to the demands of exercise.<sup>15</sup>

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## Nonstandard Abbreviations and Acronyms

<b>ACSL1</b>	acyl-coenzyme-A synthetase long-chain family member 1
<b>ActRII</b>	activin type II receptor
<b>ActRIIA</b>	activin type IIA receptor
<b>ActRIIB</b>	activin type IIB receptor
<b>AKT</b>	protein kinase B
<b>ALK</b>	activity type I receptor
<b>AMPK</b>	AMP-activated protein kinase
<b>ATF2</b>	activating transcription factor 2
<b>ATG7</b>	autophagy-related 7
<b>BAIBA</b>	$\beta$ -aminoisobutyric acid
<b>BECN1</b>	beclin-1
<b>BMP</b>	bone morphogenic protein
<b>BNP</b>	brain natriuretic peptide
<b>CAD</b>	coronary artery disease
<b>CaMKII</b>	calcium/calmodulin-dependent protein kinase
<b>CDK2</b>	cyclin-dependent kinase 2
<b>CEBP<math>\alpha</math></b>	CCAAT/enhancer-binding protein-alpha
<b>CoA</b>	coenzyme-A
<b>CPT</b>	carnitine palmitoyltransferase
<b>CREB</b>	cAMP response element-binding protein
<b>CVD</b>	cardiovascular disease
<b>eNOS</b>	endothelial nitric oxide synthase
<b>ERK</b>	extracellular signal-regulated kinase
<b>ERR<math>\alpha</math></b>	estrogen-related receptor alpha
<b>FABPpm</b>	plasma membrane-associated fatty acid binding protein
<b>FATP</b>	fatty acid transport protein
<b>FBXO32/ Atrogin1/MAFBX</b>	f-box protein 32
<b>FNDC5</b>	fibronectin type III domain-containing protein 5
<b>FOXO</b>	forkhead box O
<b>GLP1</b>	glucagon-like peptide-1
<b>GLP1RA</b>	glucagon-like peptide-1 receptor agonist
<b>GLUT4</b>	glucose transporter type 4
<b>HCAR1</b>	hydroxycarboxylic acid receptor 1
<b>HDAC</b>	histone deacetylase
<b>HF</b>	heart failure
<b>IGF</b>	insulin-like growth factor
<b>IGFBP</b>	insulin-like growth factor-binding protein
<b>IL</b>	interleukin
<b>LDL</b>	low-density lipoprotein

<b>MAPK</b>	mitogen-activated protein kinase
<b>MCT</b>	monocarboxylate transporter
<b>MEF2</b>	myocyte enhancer factor 2
<b>Metnl</b>	meteorin-like protein
<b>MFN</b>	mitofusin
<b>MOTS-c</b>	mitochondrial open reading frame of the 12S rRNA type-c
<b>mTOR</b>	mammalian target of rapamycin
<b>MuRF1/Trim63</b>	muscle ring-finger protein-1
<b>NFAT</b>	nuclear factor of activated T cells
<b>NF<math>\kappa</math>B</b>	nuclear factor kappa-B
<b>NLRP3</b>	nucleotide-binding domain, leucine-rich repeat, and pyrin domain-containing protein 3
<b>NRF1</b>	nuclear respiratory factor 1
<b>OPA1</b>	optic atrophy 1
<b>PGC1<math>\alpha</math></b>	peroxisome proliferator-activated receptor gamma co-activator 1 alpha
<b>PKA</b>	protein kinase A
<b>PPAR<math>\gamma</math></b>	peroxisome proliferator-activated receptor gamma
<b>TFEB</b>	transcription factor EB
<b>TGF<math>\beta</math></b>	transforming growth factor beta
<b>TNF<math>\alpha</math></b>	tumor necrosis factor alpha
<b>TSC2</b>	tuberous sclerosis complex 2
<b>VEGF</b>	vascular endothelial growth factor
<b><math>\beta</math>-MHC</b>	beta-myosin heavy chain

With regard to metabolic adaptations to skeletal muscle, exercise improves glucose uptake,<sup>16,17,18,19</sup> increases translocation and expression of GLUT4 (glucose transporter type 4),<sup>20,19</sup> enhances mitochondrial activity,<sup>21</sup> improves the capacity to take up and oxidize fat as fuel,<sup>21</sup> and increases the release of exercise-induced myokines into the bloodstream.<sup>22,23,24,25,26,27,3</sup> These exercise-induced adaptations improve overall metabolic and cardiovascular function.

Exerkines refer to molecules secreted in response to exercise and play a significant role in regulating various bodily functions. These molecules include proteins, metabolites, and noncoding nucleic acids, are secreted by muscles (myokines) or other organs, and can act on the organ itself (autocrine), nearby cells (paracrine), or distant organs (endocrine).<sup>28,29,30</sup> When acting in an endocrine manner, exerkines facilitate communication between different tissues and organs, including the heart, skeletal muscle, liver, and adipose tissue. By promoting this crosstalk, exerkines likely work together to enhance overall metabolic health.<sup>31</sup> Importantly, recent data indicate that endurance training induces molecular

adaptations across 19 different tissues,<sup>32</sup> many associated with mitochondrial or metabolic function.<sup>33</sup> These signaling molecules are of great interest as potential therapeutic targets due to their role in mediating the system-wide benefits of exercise. While initial research focused on myokines, exercise-induced secreted molecules can be produced by various tissues.<sup>31</sup>

In this review, we will focus on the role of skeletal muscle–released myokines in exercise-induced interorgan crosstalk with a specific focus on its role in aging and obesity.

## EXERCISE

Exercise refers to intentional physical activity and spans aerobic, resistance, high-intensity interval training, and exercise snacks.<sup>31,34,35,36</sup> Aerobic exercise is the continuous use of large muscle groups, resulting in increased muscle oxygen demand and, therefore, an increase in heart rate,<sup>37</sup> and includes activities such as walking, running, bicycling, and swimming. Resistance training is when resistance, either provided by an external source, such as weights, or by using one's body weight, is used to create progressive overload to the muscles.<sup>38</sup> High-intensity interval training is when repeated bouts of moderate-vigorous work are performed with periods of recovery of easier work or rest in between bouts.<sup>39</sup> Exercise snacks are isolated ( $\leq 1$  minute) bouts of intense exercise performed periodically throughout one's day.<sup>40</sup>

Repeated exercise, that is, exercise training, confers numerous beneficial effects on multiple tissues, including the skeletal muscle, liver, heart, vasculature, lungs, and adipose tissue.<sup>32,41</sup> Importantly, exercise improves metabolic health independent of weight loss and abrogates age-related changes in glucose disposal and insulin sensitivity.<sup>42,43,44</sup> Exercise is a potent mediator of health, and exercise capacity is the strongest predictor of mortality in humans.<sup>45,46</sup> This is achieved through the complex interplay between various signaling mechanisms that are activated in response to the exercise-induced disruption of homeostasis, ultimately leading to, for example, improved insulin sensitivity,<sup>11,47</sup> anti-inflammatory effects,<sup>48</sup> and improved cardiorespiratory fitness.<sup>15</sup>

## EXERCISE AND CARDIOVASCULAR HEALTH

CVD remains the leading cause of morbidity and mortality worldwide,<sup>49</sup> encompassing a spectrum of conditions such as arrhythmias, cardiomyopathies, heart failure (HF), and atherosclerosis.<sup>50</sup> These conditions often culminate in severe pathologies such as stroke, myocardial infarction, or cardiac arrest. The global rise in obesity has

significantly contributed to the prevalence of obesity-related CVD,<sup>51</sup> primarily through mechanisms such as hypertension and nutrient overload.<sup>52,53,54,55,56</sup> Obesity-induced hypertension promotes pathological cardiac hypertrophy, which can progress to HF.<sup>57,58</sup> In addition, the increased fatty acid uptake and utilization characteristic of an obesogenic state leads to intramyocardial lipid accumulation, lipotoxicity, and subsequent cardiac dysfunction.<sup>52,59,60</sup> Atherosclerosis, the most common form of CVD, develops gradually due to sedentary lifestyles and obesity and remains a major precursor to fatal cardiovascular events.<sup>50,61,62</sup>

Exercise training offers a powerful means to mitigate the risk and progression of CVD through multiple physiological and metabolic pathways. Regular physical activity decreases cardiovascular risk factors, including obesity, type 2 diabetes, and hypertension,<sup>63,64,65,66</sup> while improving glucose homeostasis, high-density lipoprotein levels, and blood pressure regulation, even without significant weight loss.<sup>67,68,69,70,71</sup> For patients with existing CVD, exercise-based cardiac rehabilitation enhances cardiovascular function and improves exercise tolerance,<sup>72,73</sup> which is critical in conditions such as HF.<sup>74,75,76,77,78,79</sup> Regular exercise induces favorable cardiac and vascular adaptations, including reduced resting heart rate and blood pressure, increased physiological cardiac hypertrophy, and lower circulating lipid levels.<sup>80,81,82,83</sup> Mechanistically, exercise promotes vasodilation, angiogenesis,<sup>84,85,86</sup> and the release of myokines, which mediate anti-inflammatory effects, promote exercise-induced cardiac adaptations, and facilitate intertissue communication,<sup>87,88,89,90,27</sup> fostering cardiovascular health and contributing to enhanced cardiovascular resilience.

## EXERCISE-INDUCED ADAPTATIONS TO SKELETAL MUSCLE

Cellular adaptations to skeletal muscle emerge from a complex network of signaling mechanisms triggered by exercise-induced physiological challenges.<sup>1</sup> Various stimuli, including mechanical forces such as shear stress, metabolic shifts in oxygen availability, energy substrate fluctuations, alterations in cellular pH and calcium levels, and temperature changes, collectively initiate a cascade of molecular responses.<sup>85</sup> These intricate cellular perturbations ultimately converge to modulate gene expression patterns and protein dynamics, resulting in enhanced muscular function.

### Skeletal Muscle Structural Remodeling With Exercise

Regular exercise induces structural changes in muscles to adapt to the repeated metabolic and mechanical

stresses. When muscles contract, calcium waves are generated within the myofibers due to nerve stimulation and depolarization, resulting in the activation of several signaling pathways that regulate myosin content, mitochondrial function, and muscle capillary growth. The type, intensity, and duration of calcium spikes during exercise are critical for activating calcineurin, a calcium-dependent phosphatase. Endurance exercise maximally activates calcineurin, which dephosphorylates NFAT (nuclear factor of activated T cells) transcription factors, enabling their movement into the nucleus. There, NFATs promote slow myosin expression, triggering muscle fiber-type switching to fatigue-resistant fibers.<sup>91,92</sup>

Calcineurin-NFAT signaling does not induce muscle hypertrophy, which is primarily regulated by the IGF (insulin-like growth factor)-1-AKT (protein kinase B)-mTOR (mammalian target of rapamycin) pathway. However, calcineurin activates TFEB (transcription factor EB), promoting mitochondrial biogenesis and GLUT4 expression. This enhances glucose uptake and ATP production, and supports myofiber contraction.<sup>93,94</sup> TFEB activation sustains  $\beta$ -oxidative metabolism and is partially dependent on PGC1 $\alpha$  (PPAR $\gamma$  [peroxisome proliferator-activated receptor gamma] co-activator 1 alpha), a transcription factor that promotes mitochondrial oxidative metabolism,<sup>95</sup> which is upregulated during physical activity. Interestingly, overexpression of TFEB in PGC1 $\alpha$  knockout mice can still induce mitochondrial biogenesis and improve exercise performance.<sup>94</sup> PGC1 $\alpha$  itself drives several endurance-related changes, such as mitochondrial biogenesis, fiber-type switching, fatty acid oxidation, angiogenesis, and resistance to muscle atrophy.<sup>9,96</sup>

Calcium waves also activate CaMKII (calcium/calmodulin-dependent protein kinase), which, along with calcineurin, converges on cAMP-dependent proteins such as CREB (cAMP response element-binding protein) and ATF2 (activating transcription factor 2). These proteins, when phosphorylated, bind to the promoter of the *PGC1 $\alpha$*  gene, inducing its expression. PGC1 $\alpha$  activation involves both transcriptional upregulation and posttranslational modifications that regulate protein levels and interactions with coregulators.<sup>97</sup> PGC1 $\alpha$  also plays a role in enhancing blood vessel growth by coactivating ERR $\alpha$  (estrogen-related receptor alpha), which stimulates VEGF (vascular endothelial growth factor) expression.<sup>98</sup> Endurance exercise also activates proteolytic systems such as autophagy and the ubiquitin-proteasome pathway to remodel proteins and remove damaged organelles, including mitochondria through mitophagy. This process helps prevent oxidative stress and is mediated by the energy stress sensor, AMPK (AMP-activated protein kinase).<sup>99,100,101</sup> Autophagy activation is critical for training adaptations and improved performance.<sup>102</sup>

In contrast, resistance training induces a specific variant of PGC1 $\alpha$ , known as PGC1 $\alpha$ 4, which promotes muscle hypertrophy.<sup>103</sup> PGC1 $\alpha$ 4, expressed

via an alternative promoter, does not shift muscles to oxidative metabolism but, instead, enhances protein synthesis and blocks myostatin production. The IGF1-AKT-mTOR pathway, which controls protein synthesis and muscle growth, is crucial for resistance training-induced hypertrophy and prevents muscle wasting in catabolic conditions.<sup>104,105,106,107</sup> The increases in hypertrophy as a result of myostatin inhibition on hypertrophy are partly due to its interaction with the AKT-mTOR signaling pathway.

## Exercise and Skeletal Muscle Metabolism

Myocytes undergo dynamic changes in metabolism to support both the mass and constant use of skeletal muscle. During periods of fasting and when at rest, the energetic needs are met by the oxidation of fatty acids.<sup>108</sup> Physical activity requires alterations in metabolism to support changes in membrane excitability, calcium handling, myofilament cycling, and other ATP-demanding processes. Exercise alters muscle metabolism, increasing glycogen storage, mitochondrial biogenesis, and  $\beta$ -oxidation.<sup>109,110,111,112</sup> Muscle contractions elevate cytosolic calcium levels, activating CaMKII signaling pathways, promoting glucose uptake,<sup>19,113</sup> and adrenaline-driven glycogenolysis to support energy needs.<sup>114</sup>

At rest, fatty acids in circulation are incorporated into the intramyocellular triglyceride stores before mitochondrial oxidation.<sup>115,116</sup> During exercise, fatty acids from both adipose tissue and the intramyocellular triglyceride stores are oxidized by the muscle.<sup>117–120</sup> These fatty acids are taken up by the skeletal muscle likely via fatty acid transporters such as CD36 (cluster of differentiation 36), FABPpm (plasma membrane-associated fatty acid binding protein), FATP (fatty acid transport protein)-1, and FATP4.<sup>121–124</sup> The mechanism behind increased fatty acid transporter translocation during exercise has not been elucidated but is likely, in part, regulated by AMPK<sup>125</sup> although regulation by other signaling molecules has also been suggested.<sup>126,127</sup> After entering the muscle, fatty acids are modified by ACSL1 (acyl-coenzyme-A synthetase long-chain family member 1), the key isoform in the muscle<sup>128</sup> to their acyl-CoA (coenzyme-A) form. To cross the inner mitochondrial membrane, the fatty acyl-CoAs are then further modified by CPT (carnitine palmitoyl transferase)-1 and CPT2 as part of the carnitine shuttle, after which they are oxidized.<sup>127</sup>

Exercise modulates muscle metabolism before, during, and after physical activity. Resting metabolism is altered by exercise training, increasing glycogen muscle content, mitochondrial biogenesis, and promoting  $\beta$ -oxidation.<sup>109,110,111,112</sup> In general, various exercise stimuli alter metabolism to support physical activity and promote health. Muscle contraction increases cytosolic calcium

levels, resulting in CaMKII activation and promoting glucose uptake.<sup>19,113</sup> Adrenaline stimulates glycogenolysis, promoting glycogen oxidation, and activating pyruvate dehydrogenase to promote carbohydrate oxidation.<sup>114</sup>

### Exercise Modalities Induce Specific Adaptations to Skeletal Muscle Metabolism

Exercise modality, duration, and intensity contribute to the metabolic response in the skeletal muscle. During intense exercise, the energetic requirements of skeletal muscle increase dramatically, and ATP consumption can increase as much as 100-fold. To support augmented metabolic demand, muscles initially leverage rapid energy sources, including phosphocreatine reserves and anaerobic glycogen breakdown. These initial energy-generating pathways are dynamically regulated by rapid fluctuations in key molecules such as AMP (adenosine monophosphate), ADP (adenosine diphosphate), inorganic phosphate, NAD<sup>+</sup>/NADH (nicotinamide adenine dinucleotide) ratios, and intracellular calcium concentrations released from the sarcoplasmic reticulum. Within a brief timeframe, a transition to aerobic metabolism and mobilization of additional molecular energy reserves is necessary.<sup>129,1</sup> Throwing, jumping, and sprinting are examples of short, high-intensity exertions, where ATP is generated primarily by anaerobic metabolism, with anaerobic glycolysis producing lactate,<sup>130,131</sup> which can be secreted, oxidized, and used for gluconeogenesis and muscle glycogenesis.<sup>132,133,134</sup> Skeletal muscle lactate production also regulates cardiovascular and pulmonary function during exercise. Lactate promotes angiogenesis through VEGF,<sup>135</sup> provides the metabolic substrate for cardiac metabolism,<sup>136</sup> and may act as a hypoxia sensor, upregulating breathing during exercise.<sup>137</sup>

Sustained aerobic exercise is fueled by mitochondrial oxidative phosphorylation.<sup>138,130,139</sup> The substrates for oxidative phosphorylation are primarily derived from intramuscular glycogen stores, circulating glucose, and fatty acids from within the muscle and those released into circulation by the adipose tissue.<sup>140,141</sup>

Exercise intensity dictates primary metabolic pathways, with high-intensity activities predominantly relying on carbohydrate metabolism, while moderate-intensity exercise utilizes a mix of carbohydrates and free fatty acids.<sup>141</sup> Secreted lactate can be converted to glucose or metabolized by the gut microbiota *Veillonella*, which increases in abundance in response to exercise. Higher levels of *Veillonella* in the gut result in improved exercise capacity in mice, demonstrating the intricate, cross-tissue regulation of metabolism during exercise.<sup>142</sup>

Metabolic adaptations to skeletal muscle are driven by exercise-induced changes in gene expression, protein synthesis, protein activation/inhibition, and release of signaling molecules. Exercise and training lead to extensive remodeling of the skeletal muscle epigenome,<sup>143,144</sup>

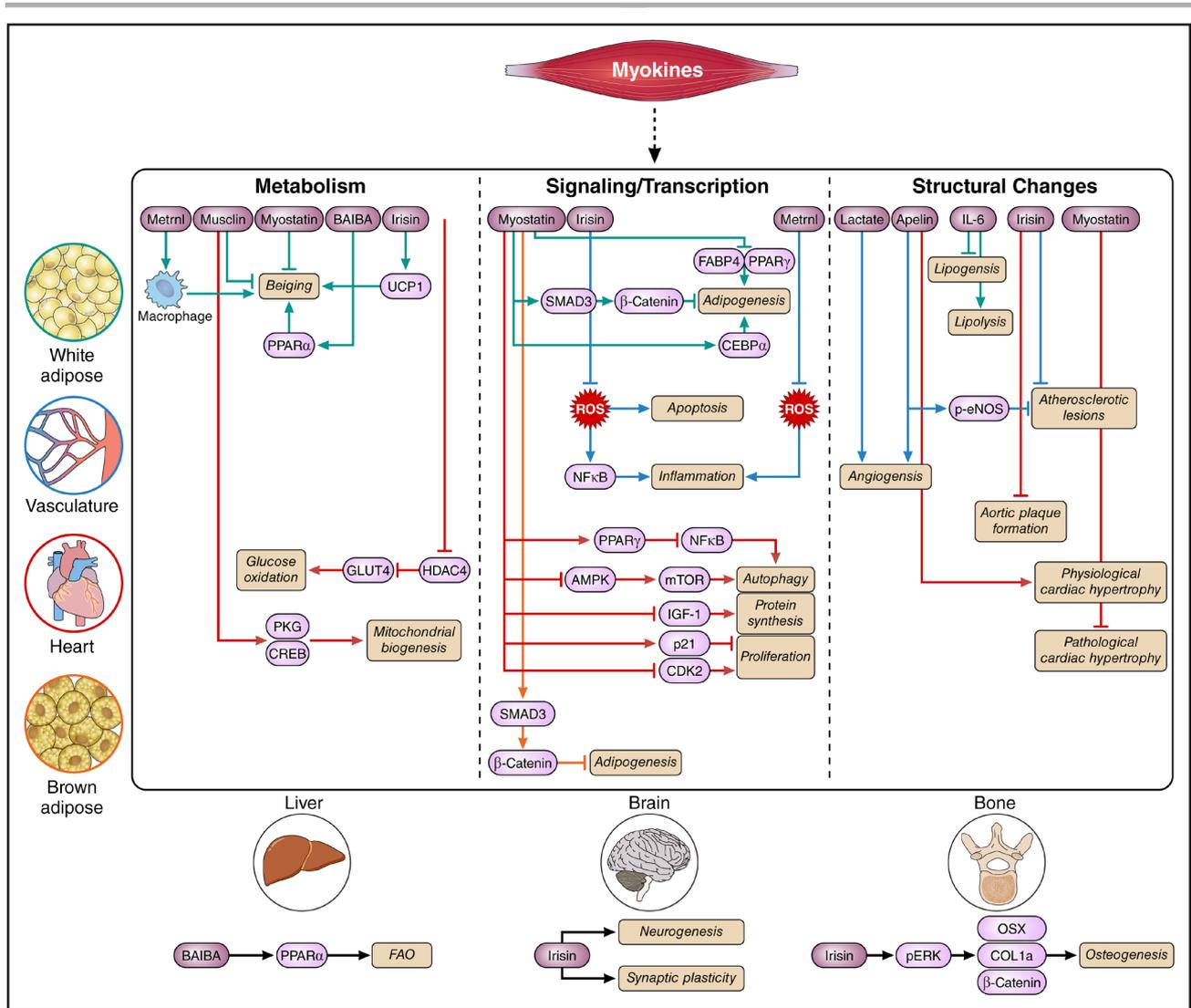
transcriptomes,<sup>145,146</sup> proteome<sup>147,148,149</sup> and various posttranslational modifications including the phosphoproteome and acetylome.<sup>148,32</sup> Critical mediators of these changes include AMPK, which is activated in skeletal muscle when ATP levels decrease, resulting in suppression of several anabolic pathways and stimulation of catabolic processes, for example, glycogenolysis. AMPK-regulated transcription factors include MEF2 (myocyte enhancer factor 2) and NRF1 (nuclear respiratory factor 1) that are important regulators of mitochondrial biogenesis. CaMKII promotes the activity of CREB, MEF2, and HDACs (histone deacetylases).<sup>1</sup> MAPKs (mitogen-activated protein kinases), mTOR, and PKA (protein kinase A) are other important mediators of the metabolic responses to exercise.<sup>129</sup> Several of these factors also increase the nuclear abundance of the cofactor PGC1 $\alpha$ .<sup>1,21</sup> The molecular drivers of the metabolic effects of exercise have been reviewed in detail elsewhere.<sup>129,1,150,127</sup> Importantly, while these central regulators are critical for exercise adaptation in skeletal muscle, the phenotypic and molecular responses demonstrate significant variability, and there is a need for further investigation into how these intricate signaling mechanisms mediate long-term metabolic health benefits of exercise training in various populations.

### SKELETAL MUSCLE FACTORS SECRETED DURING EXERCISE AND INTERORGAN CROSSTALK

Skeletal muscle releases a variety of signaling molecules, including myokines and metabolites, in response to exercise (Figure 1; Table). The term myokine was introduced in 2003<sup>185</sup>, shortly after the cytokine, IL (interleukin)-6, was identified as a skeletal muscle product released in response to exercise.<sup>186</sup> Myokines are defined as cytokines and other peptides released by the muscle, which exert various effects on the skeletal muscle itself and enter the bloodstream to regulate the function and metabolism of other organs.<sup>187,188, 29,111,30</sup> These secreted factors are differentially regulated by level of activity and intensity, as well as disease state,<sup>189,190,191</sup> with both positive and negative effects on systemic metabolism and cardiovascular health.

#### Myostatin

The myokine myostatin is released from myocytes after acute bouts of exercise<sup>192</sup> and acts within the muscle tissue. Myostatin impairs satellite cell entry into the cell cycle and protein synthesis and alters myoblast cell cycle progression, resulting in reduced muscle size.<sup>193,194</sup> Consistent exercise reduces myostatin levels, and genetic deletion or pharmacological inhibition of myostatin enhances skeletal muscle hypertrophy in



**Figure 1. Skeletal muscle releases myokines, cytokines, and metabolites in response to exercise.**

These molecules act on other organs and organ systems, such as the white adipose, vasculature, heart, brown adipose, liver, brain, and bone, to regulate changes in metabolism, signaling, transcription, and structure. Illustration credit: Sceyence Studios. AMPK indicates AMP-activated protein kinase; BAIBA,  $\beta$ -aminoisobutyric acid; CDK2, cyclin-dependent kinase 2; CEBP $\alpha$ , CCAAT/enhancer-binding protein- $\alpha$ ; Col1a, collagen type I alpha 1 chain; CREB, cAMP response element-binding protein; eNOS, endothelial nitric oxide synthase; FABP4, fatty acid binding protein 4; FAO, fatty acid oxidation; GLUT4, glucose transporter type 4; HDAC4, histone deacetylase 4; IGF-1, insulin-like growth factor; IL, interleukin; mTOR, mammalian target of rapamycin; NF $\kappa$ B, nuclear factor kappa-B; OSX, osterix; pERK, phospho extracellular signal-regulated kinase; PKG, protein kinase g; PPAR $\gamma$ , peroxisome proliferator-activated receptor gamma; p21, wildtype activating factor-1/cyclin-dependent kinase inhibitory protein-1; ROS, reactive oxygen species; SMAD3, Mothers against decapentaplegic homolog 3; UCP1, uncoupling protein 1; and 1 PPAR $\alpha$ , peroxisome proliferator activated receptor alpha.

mice<sup>195</sup> and humans,<sup>196</sup> demonstrating that myostatin is a negative regulator of exercise-induced skeletal muscle hypertrophy.

Myostatin is part of the TGF $\beta$  (transforming growth factor beta) superfamily and binds to ActRIIB (activin type IIB receptor), ActRIIA (activin type IIA receptor), and TGF $\beta$ -receptor II, activating ALK (activity type I receptor; 4, 7, 5) to phosphorylate SMAD2/3. This promotes the formation of a complex with SMAD4, leading to nuclear translocation and gene expression regulation.<sup>194</sup> Inhibition of SMAD2/3 promotes muscle growth, targeting genes involved in protein turnover. Myostatin-SMAD2/3 signaling inhibits the anabolic insulin-AKT-mTOR pathway and activates FoxO

(forkhead box O) transcription factors, increasing expression of muscle-specific E3 ligases MuRF1/Trim63 (muscle ring-finger protein-1) and FBXO32/Atrogin1/MAFBX (f-box protein 32) and inhibiting protein synthesis, leading to muscle atrophy.<sup>197</sup> The BMP (bone morphogenetic protein) signaling pathway also converges on SMAD4 to control muscle mass, with BMP-growth differentiation factor members binding to ActRIIs (activin type II receptors) and activating SMAD1/5/8 in conjunction with SMAD4. Overexpression of the BMP antagonist noggin in myostatin knockout mice suggests genetic interaction between activin-myostatin and BMP pathways. Follistatin-mediated hypertrophy blocks myostatin signaling while stimulating

**Table. Effects of myokines on target tissues.**

	BAIBA	Irisin	Metnrl	Musclin	Myostatin	Lactate	Apelin	IL-6
White adipose	Promotes beiging			Inhibits beiging	Inhibits beiging and inhibits/promotes adipogenesis			Inhibits lipogenesis and promotes lipolysis
	Roberts et al, 2014 <sup>26</sup>	Bostrom et al, 2012, <sup>22</sup> Dong et al, 2016, <sup>23</sup> Wu et al, 2012, <sup>151</sup> and Zhang et al, 2014 <sup>152</sup>	Nguyen et al, 2011 <sup>153</sup> and Rao et al, 2014 <sup>25</sup>	Jin et al, 2023 <sup>154</sup>	Artaza et al, 2005, <sup>155</sup> Arataza et al, 2005, Feldman et al, 2006, <sup>156</sup> Guo et al, 2008, <sup>157</sup> Jackson et al, 2012, <sup>158</sup> Kim et al, 2001, <sup>159</sup> Kim et al, 2012, <sup>160</sup> McPherron and Lee, 2002, <sup>161</sup> Rebbapragada et al, 2003, <sup>162</sup> and Shan et al, 2013 <sup>163</sup>			Petersen et al, 2005, <sup>164</sup> and Wan et al, 2010 <sup>165</sup>
Vasculature		Decreases apoptosis and inflammation, and inhibits atherosclerotic lesions	Decreases inflammation			Promotes angiogenesis	Promotes angiogenesis and inhibits atherosclerotic lesions	
		Lu et al, 2015, <sup>166</sup> Zhang et al, 2016, <sup>167</sup> and Zhang et al, 2016 <sup>168</sup>	El-Ashmawy et al, 2019, <sup>169</sup> Javaid et al, 2021, <sup>170</sup> and Liu et al, 2019 <sup>171</sup>			Hunt et al, 2008 <sup>135</sup>	Helker et al, 2020, <sup>172</sup> and Ishida et al, 2004 <sup>173</sup>	
Heart		Inhibits aortic plaque formation	Promotes glucose oxidation	Promotes mitochondrial biogenesis	Inhibits autophagy, protein synthesis, and proliferation, and inhibits pathological cardiac hypertrophy		Promotes physiological cardiac hypertrophy	
		Libby, 2002, <sup>174</sup> and Pober et al, 2009 <sup>175</sup>	Wang et al, 2024 <sup>176</sup>	Harris et al 2023 <sup>95</sup>	Cao et al, 2011, <sup>177</sup> Kamanga-Sollo et al, 2005, <sup>178</sup> Kamanga-Sollo et al, 2003, <sup>179</sup> and Qi et al, 2020 <sup>180</sup>		Kilpiö et al, 2024 <sup>181</sup>	
Brown adipose					Inhibits adipogenesis			
					Shan et al, 2013 <sup>163</sup>			
Liver	Promotes fatty acid oxidation							
	Roberts et al, 2014							
Brain		Promotes neurogenesis and synaptic plasticity						
		Choi et al, 2018, <sup>182</sup> and Lourenco et al, 2019 <sup>183</sup>						
Bone		Promotes osteogenesis						
		Colaiani et al, 2017 <sup>184</sup>						

BAIBA indicates β-aminoisobutyric acid; IL, interleukin; and Metnrl, meteorin-like protein.

SMAD1/5/8 activation.<sup>198,199,200</sup> Therefore, inhibiting myostatin/activins reduces phosphorylated SMAD2/3, allowing SMAD4 to interact with SMAD1/5/8 and promote

muscle growth or counteract atrophy. The BMP pathway helps prevent excessive muscle atrophy by repressing the E3 ligase FBXO30.<sup>200</sup>

Given that myostatin is also expressed in the heart,<sup>201</sup> artificially elevated levels in vivo and in vitro provide insight into the role of myostatin on cardiac function. Like the observations in skeletal muscle, myostatin treatment attenuates cardiac pathological hypertrophy in rats and isolated cardiomyocytes<sup>180</sup> (Figure 1; Table). Myostatin alters the levels of proteins involved in autophagy, a process important for maintaining appropriate cardiac function, but, in excess, autophagy is linked to the development of cardiac hypertrophy.<sup>177</sup> Myostatin downregulates AMPK-mTOR signaling and increases PPAR $\gamma$ , which leads to the silencing of NF $\kappa$ B (nuclear factor kappa-B),<sup>180</sup> a key mediator of autophagy.<sup>202</sup> Myostatin, thus, plays a crucial role in regulating cardiac hypertrophy, exerting significant effects in both in vivo and ex vivo models. Knockout or silencing of myostatin increased cardiac hypertrophy in a rat model and rat cardiomyocytes. This increase in hypertrophy is associated with an increase in key markers, such as BNP (brain natriuretic peptide) and  $\beta$ -MHC (beta-myosin heavy chain), and autophagy markers, such as LC3-II (microtubule-associated protein 1 light chain 3-II) and BECN1 (beclin-1). Furthermore, myostatin blocks cardiomyocyte proliferation by increasing P21 expression, which inhibits cell cycle progression, and reducing CDK2 (cyclin-dependent kinase 2), a critical regulator of the cell cycle. Inhibition of CDK2 availability results in cell cycle arrest. Myostatin also inhibits the proliferation of porcine embryonic myogenic cells, in part through the production of IGFBP (IGF-binding protein)-3 and IGFBP-5, which sequester IGFI outside the cell and may impede the intracellular and nuclear actions required for growth.<sup>203,202,178,179,204,180,205</sup>

Myostatin also regulates adipose tissue function through different mechanisms in vivo and in vitro (Figure 1; Table). In vitro, myostatin inhibits adipogenesis in 3T3-L1 and human mesenchymal stem cells and primary brown adipocytes through the activation of SMAD3 and  $\beta$ -catenin<sup>157,159,160,162</sup> but promotes adipogenic commitment while impairing differentiation in C3H10T1/2 cells.<sup>156</sup> However, when applied after differentiation induction, myostatin enhances adipogenesis,<sup>155</sup> suggesting stage-specific effects. In vivo, myostatin knockout mice show reduced adiposity,<sup>158,161</sup> with decreased expression of adipogenesis markers, CEBP $\alpha$  (CCAAT/enhancer-binding protein-alpha), and PPAR $\gamma$ , indicating impaired adipogenesis. However, this is likely due to glucose diversion from adipose tissue rather than direct effects on adipocyte turnover.<sup>206,207</sup> In high-fat diet-fed mice, adipose-specific loss of myostatin signaling does not significantly affect lean or fat mass, glucose, insulin, or adipokine levels.<sup>206</sup> While myostatin knockout enhances the being of subcutaneous white adipose tissue,<sup>163</sup> recombinant myostatin has no direct impact on lipid release or adipose tissue mass, highlighting the context-dependent nature of myostatin's role in adipose regulation.

## Irisin

Irisin is a myokine induced during exercise in mice and humans<sup>22,208,209,210</sup> that is secreted primarily from the skeletal muscle.<sup>211</sup> Irisin may regulate beneficial adaptations of exercise, such as improved energy expenditure,<sup>212</sup> glucose homeostasis,<sup>22</sup> and bone health.<sup>213</sup> Acute treatment of irisin on myocytes in vitro increased glucose uptake and glycolysis, and longer treatment induced mitochondrial biogenesis and increased oxygen consumption, suggesting that irisin promotes oxidative metabolism in myocytes.<sup>214</sup> Moreover, irisin-mediated changes in skeletal muscle metabolism may aid in maintaining skeletal muscle integrity in aging.<sup>189</sup> Increased muscle irisin secretion in mice improved glucose tolerance and lowered fasting insulin levels,<sup>22</sup> and levels are decreased in patients with type 2 diabetes. Circulating irisin levels are positively correlated with bone mechanical properties in humans<sup>213</sup> and induce the expression of genes associated with bone formation, such as *Osx*, *Col1a*, and *Ctnnb1*, preventing bone mineral density loss from disuse in mice.<sup>184</sup>

The relationship between irisin and cardiac health remains unclear, with conflicting data in the literature. Some studies report a positive association between serum irisin concentration and CVD, atherosclerosis, and stroke,<sup>215,216,217</sup> while others indicate negative associations and potential cardioprotective effects. For example, several studies have found lower irisin levels in patients with coronary artery disease (CAD).<sup>218,219,220</sup> In contrast, in mouse models of atherosclerosis, irisin treatment reduces the development of carotid and aortic plaques,<sup>166,167,168</sup> indicating a protective role (Figure 1; Table). Irisin also reduced disease severity in both genetic and surgical models of atherosclerosis.<sup>167</sup> In models of atherogenesis, oxidized LDL (low-density lipoprotein) promotes inflammation in endothelial cells, leading to apoptosis and plaque formation.<sup>174,175</sup> In vitro, irisin treatment attenuated oxidized LDL-induced inflammation and apoptosis in endothelial cells by decreasing reactive oxygen species and suppressing inflammation and apoptosis via eNOS (endothelial nitric oxide synthase) phosphorylation and decreased NF $\kappa$ B signaling (Figure 1; Table).<sup>167</sup> These conflicting findings highlight the need for further research to clarify the mechanistic relationship between irisin and CVD.

Irisin may promote the remodeling of adipose tissue by stimulating the browning of white adipocytes and thermogenesis (Figure 1; Table). Acting directly on adipocytes, irisin enhances their thermogenic capacity by activating p38 MAPK and ERK (extracellular signal-regulated kinase) signaling pathways.<sup>152</sup> This molecular signaling cascade facilitates the conversion of white adipocytes into beige, thermogenically active cells.<sup>22,23,151,152</sup> The effects of irisin, stimulating thermogenesis and adipocyte browning, position it as a key factor in modulating

adipose tissue function and combating metabolic disorders.

Exercise has a profound impact on brain health, improving cognitive function and plasticity, particularly in older adults.<sup>221,222</sup> Regular physical activity enhances outcomes in neurodegenerative diseases and stroke.<sup>223,224,225,226</sup> Exercise also promotes neurogenesis in the adult brain, enhancing synaptic plasticity and spatial learning.<sup>227,228</sup> Irisin has been linked to improved cognitive function in various populations. Higher irisin levels correlate with better cognitive performance in older adults at risk for dementia<sup>229</sup> and young athletes.<sup>230</sup> However, elevated irisin levels in obese women have been associated with poorer executive function,<sup>231</sup> suggesting a complex relationship between irisin, cognition, and metabolic health.

Irisin has neuroprotective effects in conditions such as Alzheimer disease, Parkinson disease, stroke, and diabetes. In Alzheimer disease, irisin supports hippocampal neurogenesis and gene regulation,<sup>182</sup> improving memory and synaptic plasticity in mouse models (Figure 1; Table).<sup>183</sup> FNDC5 (fibronectin type III domain-containing protein 5), the precursor to irisin, also enhances memory and synaptic function in Alzheimer disease. In female Alzheimer disease mice, irisin reduced tau protein levels and inflammation although it worsened inflammation in males.<sup>232</sup> In addition, irisin reduces amyloid-beta-induced inflammation in astrocytes and promotes hippocampal cell proliferation.<sup>233</sup>

In Parkinson disease, irisin preserves dopaminergic neurons, improves motor function, and aids stem cell migration and differentiation in rat models.<sup>234</sup> Following ischemic stroke, where irisin levels decline, increasing irisin reduces brain infarct size, inflammation, neurological deficits, and brain edema in mouse models.<sup>235</sup> In vitro irisin reduced oxidative stress and inflammation in neurons under ischemic conditions.<sup>236</sup> In diabetic mouse models, irisin improved cognitive function, preserved synaptic proteins, reduced inflammation, and attenuated NF $\kappa$ B activation in the brain, suggesting its potential to counteract diabetes-associated neurodegeneration.<sup>237</sup> Collectively, these studies highlight the broad interorgan signaling effects of a myokine and demonstrate the therapeutic potential of irisin for various neurological and metabolic disorders.

### Meteorin-Like 1

The myokine Metrnl (meteorin-like protein) was identified in primary myotubes overexpressing the PGC1 $\alpha$  splice isoform, PGC1 $\alpha$ 4, which regulates muscle growth and energy expenditure.<sup>103</sup> Metrnl is induced in the skeletal muscle after exercise or cold exposure,<sup>25</sup> suppressing inflammation by inhibiting inflammasome activation.<sup>170</sup> Exogenous Metrnl administration in mice fed a high-fat diet protects from lipid-induced insulin resistance in skeletal muscle and C2C12 cells.<sup>238</sup> Moreover, higher levels of Metrnl in circulation increase energy expenditure and

improve glucose homeostasis in mouse models of obesity and diabetes.

Exercise is associated with decreased mortality, increased quality of life, and higher rehabilitation success in patients with CVD.<sup>239,240,241</sup> In a mouse model of HF, Metrnl administration improved cardiac structure and function, whereas Metrnl knockdown eliminated these benefits.<sup>176</sup> Metrnl activates AMPK, which suppresses histone deacetylases such as HDAC4,<sup>242,243</sup> leading to increased GLUT4 expression.<sup>244,245,246</sup> Impaired energy metabolism is a hallmark of HF, and Metrnl-induced GLUT4 expression promotes glucose metabolism, improving mitochondrial function in the hearts of a mouse model of HF (Figure 1; Table).<sup>176</sup> These findings suggest that exercise improves cardiovascular health in HF by altering cellular metabolism via Metrnl.

Exercise also reduces reactive oxygen species and inflammation though the mechanisms remain unclear.<sup>247</sup> Elevated Metrnl levels are associated with a lower risk of CAD and are negatively correlated with LDL and inflammatory cytokines, key mediators of CAD.<sup>169,248</sup> In both a patient study and a mouse model of CAD, exercise increased Metrnl levels, which were associated with reduced LDL and inflammatory cytokines.<sup>170</sup> In addition, exercise-induced Metrnl elevation suppressed oxidative stress and inhibited the NLRP3 (nucleotide-binding domain, leucine-rich repeat, and pyrin domain-containing protein 3) inflammasome in obese mice. These findings suggest that exercise-induced Metrnl may improve CAD risk by regulating lipid metabolism and reducing inflammation (Figure 1; Table).

Metrnl plays a critical role in regulating adipose tissue function and systemic energy balance. Increased circulating Metrnl enhanced whole-body energy expenditure, promoted the browning of white adipose tissue, and improved glucose tolerance in obese and diabetic mice.<sup>25</sup> Metrnl does not act directly on adipocytes; instead, it exerts its prothermogenic effects through immune cell modulation. Metrnl recruits eosinophils to adipose, which release IL-4 and IL-13, driving the alternative activation of adipose tissue macrophages. These alternatively activated macrophages are key mediators of cold-induced thermogenesis (Figure 1; Table)<sup>153</sup> and exert anti-inflammatory effects, reducing adipose inflammation and contributing to metabolic health. Through immune regulation, Metrnl establishes a unique mechanism of thermogenic activation and inflammation modulation in adipose tissue.

### Mitochondrial Open Reading Frame of the 12S rRNA Type-c

MOTS-c (mitochondrial open reading frame of the 12S rRNA type-c) is a mitochondria-derived peptide that is increased with exercise in humans, expressed in the skeletal muscle,<sup>249</sup> and regulates metabolism.<sup>250,251</sup> MOTS-c targets the folate cycle and increases levels of 5-aminoimidazole-4-carboxamide-1- $\beta$ -D-ribofuranoside,

leading to the activation of AMPK.<sup>251</sup> Expression of MOTS-c is age-dependent, circulating levels are negatively correlated with HbA1c levels in patients with type 2 diabetes,<sup>252</sup> and expression of MOTS-c is lower in both skeletal muscle and serum of patients with chronic kidney disease.<sup>248</sup> Genetic variation in MOTS-c has been associated with human longevity,<sup>253</sup> and when given to mice, it reverses age-dependent insulin resistance, suggesting that it may have therapeutic potential in metabolic and aging-associated diseases.<sup>251</sup>

### $\beta$ -Aminoisobutyric Acid

$\beta$ -Aminoisobutyric acid (BAIBA) is a small molecule of myokine that has been shown to promote adipose tissue beiging and fatty acid metabolism in the liver through regulation of PPAR $\alpha$  (peroxisome proliferator-activated receptor alpha) (Figure 1; Table).<sup>26</sup> BAIBA has also been implicated in bone health through the prevention of osteocyte apoptosis.<sup>254</sup> BAIBA was identified as a potential myokine through liquid chromatography-mass spectrometry metabolic profiling of human myocytes overexpressing PGC1 $\alpha$ . When BAIBA was added to the drinking water of mice, it significantly boosted the expression of beiging-related genes, such as *Ucp1* and *Cidea*, in subcutaneous white adipose tissue.<sup>3</sup> Similarly, when primary stromal-vascular fraction cells isolated from white adipose tissue were incubated with BAIBA, *Ucp1* and *Cidea* expression was increased. Human pluripotent stem cells exposed to BAIBA during differentiation into mature white adipocytes also showed elevated expression of beiging markers. Exercise significantly elevated BAIBA levels in both mice and humans. In mice, 3 weeks of voluntary wheel-cage running led to a notable increase in circulating BAIBA, while, in humans, 20 weeks of supervised submaximal aerobic exercise training<sup>255</sup> in 80 subjects resulted in a significant rise in circulating BAIBA.<sup>256</sup> These findings suggest that exercise significantly boosts circulating BAIBA in both mice and humans, and in rodents and isolated human cells, BAIBA may play a role in promoting the beiging of subcutaneous white adipose tissue. However, the direct involvement of BAIBA in exercise-induced beiging in human subcutaneous white adipose tissue remains to be fully established.

### Musclin

Musclin, encoded by the *Ostn* gene, is secreted by skeletal muscle in response to exercise. During exercise, its production is stimulated by calcium-induced activation of AKT1, which removes transcriptional inhibition of the *Ostn* by FOXO1.<sup>257</sup> Musclin has been shown to inhibit proliferation of fibro-adipogenic progenitor cells in skeletal muscle through upregulation of filamin A interacting protein 1 like, which promotes fibro-adipogenic progenitor apoptosis, thereby reducing fibrosis and abnormal fatty infiltration.<sup>258</sup>

In mice, when *Ostn* is disrupted and musclin secretion is eliminated, exercise tolerance is diminished; administration of recombinant musclin rescues this phenotype. The reduction in exercise capacity is associated with lower plasma levels of atrial natriuretic peptide and diminished levels of cyclic guanosine monophosphate and PGC1 $\alpha$  in skeletal muscles of knockout animals following exercise.<sup>257</sup> Secreted musclin similarly promotes mitochondrial biogenesis in cardiac muscle (Figure 1; Table).<sup>95</sup> However, running capacity remains the same in mice with muscle-specific knockdown of musclin.<sup>154</sup> Paradoxically, obesity is associated with higher circulating levels of musclin. Overexpression of musclin reduces thermogenesis in beige fat (Figure 1; Table), while inactivation has the opposite effect and improves glucose tolerance and insulin sensitivity.<sup>154</sup> Importantly, transient activation of musclin shows many cross-tissue metabolic benefits, while continuous activation appears detrimental to metabolic health.

In cardiac muscle, musclin plays a critical role in promoting mitochondrial biogenesis in response to exercise (Figure 1; Table). Disrupting musclin signaling in mice eliminates exercise's ability to protect against cardiac ischemic injury. The proposed mechanism involves a musclin-induced increase in cyclic guanosine monophosphate signaling, leading to increased protein kinase G and CREB activity that together stimulate the expression of PGC1 $\alpha$ . Musclin also elevates cardiac C-type natriuretic peptide levels, which improves cardiomyocyte contractile function.<sup>259</sup> Targeted infusion of musclin reproduced the cardioprotective benefits of exercise in both sedentary wild-type and *Ostn*-knockout mice.<sup>95</sup> Similarly, skeletal muscle-specific knockout mice were more sensitive to pressure overload, and *Ostn* expression is lower in the skeletal muscle of patients with HF.<sup>259</sup>

### Apelin

Apelin is an exercise-induced myokine<sup>260</sup> that is also released from adipose tissue in response to insulin signaling.<sup>261</sup> Apelin affects several organs, including the heart, brown adipose tissue, brainstem, and kidneys.<sup>261,262,263,264,265</sup> It functions by binding to its G-protein coupled receptor, which triggers intracellular signaling that activates AMPK, promoting mitochondrial biogenesis and improving glucose uptake in skeletal muscle.<sup>266</sup>

In obese, insulin-resistant mice, apelin treatment improves insulin sensitivity.<sup>261</sup> In addition, apelin stimulates angiogenesis by activating endothelial cells and promoting glycolytic activity, such as through the activation of c-MYC,<sup>172</sup> and it helps lower arterial blood pressure, potentially via phosphorylation of eNOS (Figure 1; Table).<sup>173</sup> In apelin knockout mice, high-intensity interval training failed to induce insulin-like growth factor 1 and resulted in smaller type I muscle fibers compared with wild-type mice.<sup>181</sup>

In the heart, apelin induces physiological cardiac hypertrophy, enhances mitochondrial gene expression,

and increases ATP production (Figure 1; Table).<sup>181</sup> It also has cardioprotective effects, especially in mice prone to atrial fibrillation.<sup>263</sup> For instance, apelin knockout mice developed eccentric rather than concentric left ventricular hypertrophy in response to high-intensity interval training,<sup>181</sup> highlighting the role of exercise-induced apelin expression in promoting normal cardiac remodeling. Apelin has also been linked to the transgenerational effects of exercise on metabolic health.<sup>265</sup>

## Brain Natriuretic Peptide

BNP is primarily secreted by the heart in response to increased cardiac wall stress but is also produced by skeletal muscles. Supraphysiological BNP levels in mice protect against high-fat diet-induced obesity and insulin resistance and stimulate mitochondrial biogenesis in skeletal muscle through upregulation of PGC1 $\alpha$  and PPAR $\delta$ .<sup>267</sup> BNP also has immunomodulatory functions though the exercise-induced benefits from this modulation are unknown.<sup>268</sup>

## IL-6

IL-6, an inflammatory cytokine, is released from myocytes during exercise,<sup>24</sup> increasing in concentration with the length of the physical activity.<sup>269</sup> IL-6 is produced during the contraction of skeletal muscle in a TNF $\alpha$  (tumor necrosis factor alpha)-independent manner, uncoupling IL-6 production from the traditional inflammatory cascade.<sup>270</sup> IL-6 secretion has been reported *ex vivo* in cultured human myotubes<sup>271</sup> and activates AMPK, enhancing energy consumption.<sup>24</sup> IL-6 regulates skeletal muscle fatty acid metabolism and glucose uptake<sup>272,273</sup> and promotes whole-body fatty acid oxidation.<sup>274</sup> IL-6 release is enhanced when glycogen levels are low, suggesting that IL-6 acts as an energy sensor in skeletal muscle rather than a mediator of inflammation.<sup>275,190,276,277</sup>

IL-6 signaling is associated with adipocyte inflammation in obesity,<sup>278,274</sup> but its effects on adipose tissue can vary in different contexts. In obesity, IL-6 plays a role in metabolic programming, particularly in skeletal muscle, and may have similar effects in white adipose tissue. Global IL-6 knockout mice show higher body weight due to increased subcutaneous white adipose mass,<sup>279,280</sup> suggesting IL-6 regulates adipose tissue maintenance and metabolism. IL-6 enhances lipolysis in 3T3-L1 cells<sup>164</sup> and activates AMPK (Figure 1; Table).<sup>281</sup> In IL-6 knockout mice, exercise increases PPAR $\gamma$  expression in subcutaneous white adipose,<sup>279</sup> indicating that IL-6 may influence transcription factors involved in adipogenesis and adipocyte maintenance. However, IL-6 knockout does not alter AMPK activity in subcutaneous white adipose tissue<sup>279</sup> though it impairs exercise-induced AMPK activation in adipocytes,<sup>281</sup> suggesting that IL-6 is required for this process. IL-6

also promotes *Ucp1* expression and thermogenesis in subcutaneous white adipose tissue, contributing to the being effect of exercise.<sup>279</sup> In addition, IL-6 dampens the exercise-induced induction of gluconeogenic enzymes such as pyruvate dehydrogenase kinase 4 in white adipose,<sup>165</sup> indicating that IL-6 impairs lipogenesis during exercise.

The role of IL-6 in glucose metabolism is less clear. Some studies show that IL-6 treatment enhances glucose uptake in adipocytes,<sup>272,282</sup> while others find no effect.<sup>283,284</sup> In humans, IL-6 infusion increases glucose uptake in subcutaneous white adipose tissue<sup>285</sup> but does not promote lipolysis or activate IL-6 signaling in adipose tissue from healthy individuals.<sup>273</sup> These findings underscore the complex, context-dependent role of IL-6 in regulating adipose tissue, balancing lipid metabolism, glucose handling, and adipogenesis in response to exercise.

## IL-8

Expression of the chemokine IL-8 is increased in skeletal muscle in response to endurance and resistance exercise in humans<sup>286,287</sup> and is predominantly detected after extended physical exertion such as marathon running.<sup>288</sup> Transcription is associated with a reduction in glycogen levels<sup>289</sup> and results in antiapoptotic and proangiogenic signaling through CXCR2 receptor interactions. In endothelial cells, for example, IL-8 increases protein levels of the antiapoptotic factor B-cell lymphoma 2 and matrix metalloproteinases 2 and 9.<sup>290</sup> While IL-8 is induced to a similar extent in young and old individuals, training attenuates the acute IL-8 response in older individuals,<sup>287</sup> potentially reducing the acute, proinflammatory effects of exercise.

## Lactate

Lactate is a product of glycolysis under aerobic conditions.<sup>136</sup> Muscle glucose uptake is increased with exercise,<sup>291</sup> increasing lactate production and secretion.<sup>136</sup> Elevated generation of lactate in muscles during exertion regenerates NAD levels, increasing the glycolytic capacity of the cell. Lactate can also modulate skeletal muscle mitochondrial expansion and metabolism. Lactate stimulates mitochondrial biogenesis upstream of PGC1 $\alpha$ .<sup>292</sup> Moreover, glycolysis and the production of lactate modulate muscle fatty acid oxidation and uptake. Glycolysis increases the production of the metabolite malonyl-CoA, which inhibits CPT1, and lactate downregulates the expression of CPT2,<sup>136</sup> both of which are important enzymes in fatty acid oxidation.

Outside of the skeletal muscle itself, skeletal muscle lactate production also regulates cardiovascular and pulmonary function during exercise (Figure 1; Table). Lactate promotes angiogenesis through stimulation of VEGF,<sup>135</sup> provides metabolic substrate for cardiac function,<sup>136</sup> and upregulates breathing.<sup>137</sup> Secreted lactate

can also be converted into N-lactoyl-phenylalanine by various immune and endothelial cells, and it increases substantially in blood in response to one bout of exercise. Treatment with N-lactoyl-phenylalanine has been demonstrated to improve glucose metabolism and decrease adiposity in mice.<sup>293</sup>

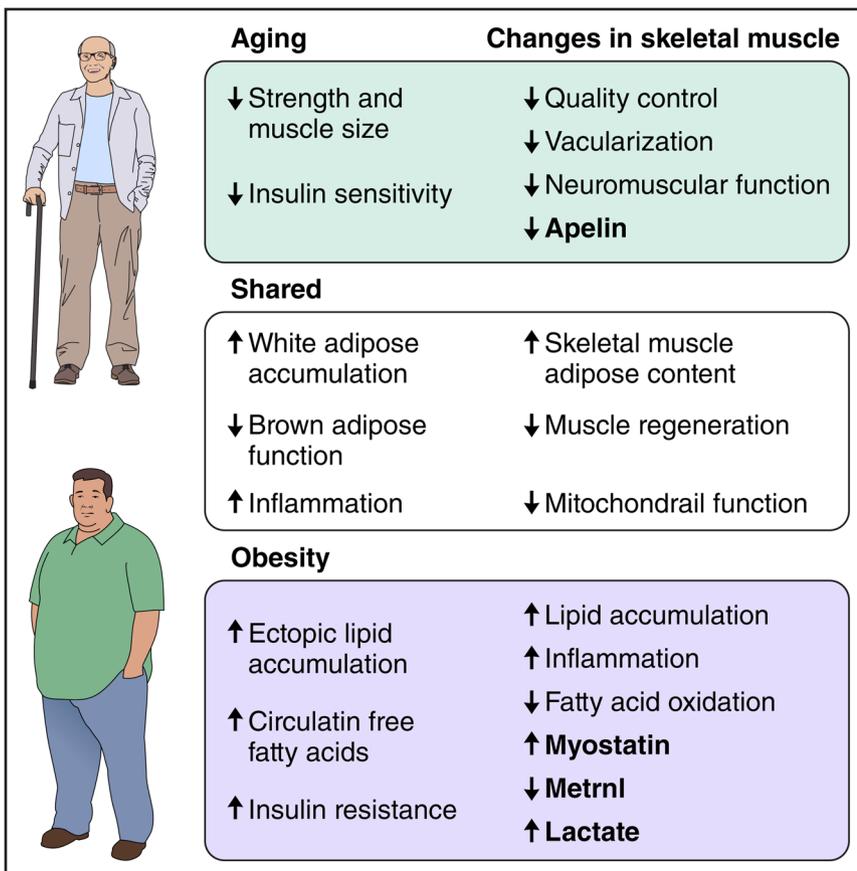
Skeletal muscle-derived lactate plays a multifaceted role in regulating adipose tissue (Figure 1; Table). High-intensity physical activity results in lactatemia and a corresponding drastic decrease in plasma-free fatty acid concentrations. This is due to the activation of the HCAR1 (hydroxycarboxylic acid receptor 1) on adipocytes by lactate, inhibiting lipolysis in both rodents and humans.<sup>136</sup> Beyond its antilipolytic effects, lactate promotes thermogenic adaptations in adipose tissue. Lactate induces the browning of white adipose, contributing to increased thermogenic capacity. This effect is mediated by proton-linked MCTs (monocarboxylate transporters) present in adipocytes, which facilitate lactate transport into cells.<sup>294</sup> Lactate increases the expression of thermogenic genes, including *Ucp1*, through pathways dependent on PPAR $\gamma$  signaling and intracellular redox modifications.<sup>295</sup> In brown adipose tissue, *Mct1* expression is significantly elevated after exercise, aligning with physiological stimuli of browning and enhancing lactate import for metabolic adaptation.<sup>296</sup> Together, these processes highlight lactate's

dual role as both a regulator of lipid mobilization and a potent inducer of thermogenic programming in adipose tissue.

## AGING AND LONGEVITY

Increased life expectancy has resulted in older adults becoming the fastest-growing subpopulation in the United States. Aging is the leading driver of disease and mortality,<sup>297</sup> and caring for the aging population presents a significant burden on the health care system.<sup>298</sup> Thus, therapeutic strategies prioritizing the prevention of age-associated decreases in mobility and metabolic decline are of utmost importance. Aging is associated with changes to many metabolic tissues, including the atrophy of skeletal muscle,<sup>299</sup> increase in visceral white adipose tissue depots,<sup>300,301,302</sup> and a decrease in the function of brown adipose tissue (Figure 2; Table).<sup>303</sup> Changes in these depots are both consequences of and drivers for age-associated metabolic dysfunction, emphasizing the importance of how aging disrupts appropriate function in multiple tissues.

Aging is a process of decline in biological function that is mostly irreversible. The causes of decline are multifactorial and driven by the accumulation of stress and damage over an organism's lifetime. Aging increases the risk of many diseases,<sup>304</sup> driven by decreased physical



**Figure 2. Aging and obesity both induce changes in white adipose accumulation, brown adipose function, and inflammation that can impair metabolism and promote disease pathogenesis.**

Obesity promotes changes in muscle lipid metabolism, inflammation, and myokine secretion, and both states increase skeletal muscle adipose content, impair muscle regeneration, and decrease mitochondrial function. Despite the similarities, less is known about how aging alters myokine release and whether the effects differ between sexes. Illustration credit: Scyeence Studios.

activity,<sup>111</sup> increased inflammation,<sup>305,111</sup> muscle atrophy, and increased central adiposity. Increased adiposity caused by energy imbalances, driving metabolic inflexibility and implicating mitochondrial dysfunction in age-associated disease progression.<sup>306,307</sup> Significantly, caloric restriction is one of the only interventions to extend lifespan, implicating systemic energy balance and metabolism in longevity.<sup>308,309</sup>

## EXERCISE AND AGING

There is an age-dependent decline in maximal heart rate, reducing the ability to increase cardiac output in response to exercise. The increase in stroke volume is also shifted from enhanced contractility to increased end-diastolic volumes in older individuals. This is due to diminished sensitivity to  $\beta$ -adrenergic signaling, impaired calcium handling and mitochondrial dysfunction, all of which can be improved with exercise training.<sup>310</sup> Exercise, thus, demonstrates promise for prevention and treatment of age-associated decline in cardiac function although further research is needed to establish optimal dose and intensity.

Advanced age is associated with structural alterations that impair muscle function (Figure 2; Table). These include increases in muscle adipose content and decreases in neuromuscular function and vascularization.<sup>311,12,312,313</sup> Aging skeletal muscle also demonstrates a decline in satellite cell number and function<sup>314,315,316</sup> that contributes to delayed<sup>317</sup> or impaired skeletal muscle regeneration after an injury.<sup>318</sup>

The dysfunction of proteostasis and organelle quality control systems plays a major role in bioenergetic defects, fatigue, and reduced force transmission in skeletal muscle as with age. A sedentary lifestyle and reduced physical activity contribute to and accelerate the onset and progression of sarcopenia. This leads to a decrease in mechanotransduction signaling, which affects protein turnover, calcium homeostasis, and mitochondrial function. The impact of reduced physical activity on sarcopenia becomes especially prominent during periods of immobilization, hospitalization, or bed rest in older individuals.<sup>319</sup>

Interestingly, inactivity suppresses mitochondrial fusion protein expression before any changes in mitochondrial mass.<sup>320</sup> However, higher expression of these proteins correlates with muscle force and bioenergetics in the elderly.<sup>321,322</sup> Inhibition of the mitochondrial fusion protein OPA1 (optic atrophy 1) in adult mice leads to premature aging features, including increased oxidative stress, systemic inflammation, sarcopenia, and senescence markers, ultimately causing death.<sup>323,322</sup> Similarly, deletion of the fusion protein MFN (mitofusin)-2 in muscle causes sarcopenia,<sup>324</sup> and double deletion of MFN1 and MFN2 results in severe muscle loss and mitochondrial DNA depletion.<sup>325</sup> On the other hand, deleting the

fission protein, dynamin-related protein 1, causes myopathy, with myofiber degeneration due to imbalanced calcium homeostasis and mitochondrial-endoplasmic reticulum tethering.<sup>326</sup>

Gain-of-function experiments further demonstrate the effects of mitochondrial shaping proteins on muscle mass. Overexpression of OPA1 counters muscle loss,<sup>327</sup> while upregulation of fission machinery promotes muscle wasting.<sup>328,329</sup> Mitochondrial dynamics, when altered, profoundly affect mitochondrial function and the onset of sarcopenia, with physical activity playing a key role. Mitochondrial dynamics are interconnected with mitophagy, mitochondrial proteostasis, and the broader mitochondrial quality control system, which adapts mitochondrial bioenergetics to muscle needs.

Inhibition of autophagy, through ATG7 (autophagy-related 7) deletion or TSC2 (tuberous sclerosis complex 2) overexpression in muscle, mimics the pathological features of age-related sarcopenia, such as increased oxidative stress, dysfunctional mitochondria accumulation, myofiber denervation, atrophy, weakness, and premature death.<sup>330,331</sup> Reactivating autophagy in aged mice, either by expressing ATG7 in skeletal muscle or treating with urolithin A (a compound that promotes mitophagy), helps restore muscle mass.<sup>330,332</sup> Overexpression of sestrins, stress-responsive proteins that decline with age, also promotes autophagy and reduces muscle wasting and weakness in older mice.<sup>333</sup> Importantly, failure of autophagy is linked to mitochondrial dysfunction, oxidation of contractile proteins, and impaired force generation.<sup>330</sup> Similarly, urolithin A has been shown to improve muscle strength and performance in middle-aged individuals by restoring genes and proteins involved in mitochondrial metabolism, mass, and PTEN (phosphatase and TENSin homolog)-induced putative kinase 1/Parkin-mediated mitophagy.<sup>334</sup> Importantly, urolithin A demonstrates similar effects in cardiac muscle, and supplementation in humans reduces ceramide levels in plasma, which is predictive of CVD risk.<sup>335</sup>

Mitochondrial dysfunction and impaired mitophagy contribute to sarcopenia through several mechanisms (Figure 2; Table), including reduced mitochondrial ADP sensitivity,<sup>336</sup> increased mitochondrial reactive oxygen species production and DNA damage,<sup>337</sup> activation of an inflammatory response,<sup>323,322</sup> decreased cytosolic calcium due to sequestration,<sup>326</sup> and metabolic crisis.<sup>338</sup> While autophagy is clearly important in sarcopenia, the role of the ubiquitin-proteasome system in this process remains less well understood.

Epigenetic clocks have emerged as a tool to investigate tissue aging and the impact of different antiaging therapies.<sup>339</sup> Endurance exercise training alters the human skeletal muscle genome,<sup>143</sup> and a skeletal muscle-specific epigenetic clock<sup>340</sup> has been used to demonstrate that exercise training can counteract age-related epigenetic changes in skeletal muscle. A comprehensive

meta-analysis of over 3000 human skeletal muscle samples demonstrated that higher aerobic fitness correlated with younger epigenetic and transcriptomic profiles, and exercise training can shift these patterns toward a younger state. Conversely, muscle disuse accelerated molecular aging.<sup>144</sup>

## AGING AND SARCOPENIA

Muscle function and metabolism are closely linked to health span and longevity.<sup>12,35</sup> Muscle mass typically begins to decline in the third or fourth decade of life, with the rate accelerating over time, leading to significant loss in later years.<sup>341,342</sup> This decline is driven by changes in muscle structure, function, and regeneration, with strength loss often outpacing size reduction, indicating a decline in muscle quality (Figure 2; Table).<sup>343,342,312,344</sup> When muscle aging results in excessive force loss, fatigue, and exercise intolerance, it is considered sarcopenia, a disease that predicts frailty, poor quality of life, and increased morbidity and mortality.<sup>345,346</sup>

Sarcopenia affects 15% of individuals at the age of 65 years and up to 50% by the age of 80 years.<sup>347,319</sup> While the mechanisms behind sarcopenia are not fully understood, genetic factors and lifestyle-related epigenetic changes play a significant role. Parabiosis experiments in mice suggest that serum factors from young blood can influence muscle cell senescence, indicating that hormonal, metabolic, and gene expression changes affect key determinants of muscle function, including calcium ions, ATP production, and contractile protein quality and assembly.<sup>348</sup>

In addition to reduced strength and mobility, sarcopenia also disrupts whole-body metabolism, contributing to impaired glycemic regulation and insulin resistance, which are common with age.<sup>349,350,351</sup>

## AGING ALTERS MYOKINES

Exercise continues to be an effective intervention for counteracting the effects of aging, delaying the development of disability and dependence in older adults.<sup>352,2,353</sup> The beneficial effects of exercise are, in part, mediated by secreted factors, including myokines. There are well-established, sex-based differences in many parameters in response to exercise<sup>354,355,356,357,358,359,360</sup> including in myokine release.<sup>361,362,363,364,365</sup> However, how aging alters myokine release, and the significance of age-associated disease, remains unclear (Figure 2; Table).<sup>366,367</sup> In young subjects, increased myostatin levels are associated with decreased muscle mass. However, in elderly men, low serum levels of myostatin are associated with lowered skeletal muscle mass but not in older women.<sup>367</sup> In response to endurance training, older individuals demonstrate a greater skeletal muscle

induction of interferon-induced inflammatory genes,<sup>145</sup> while single-cell transcriptional profiling showed exercise reversed aging-induced inflammatory changes in many skeletal muscle cell types.<sup>368</sup> In addition, little work has been done, which examines changes in the myokine profile in postmenopausal women. On average, women live longer than men but experience higher rates of disability and have an increased risk of age-related comorbidities that cannot be attributed solely to life span.<sup>369,370</sup>

## OBESITY

Obesity is a disease characterized by increased pathological adiposity and ectopic lipid accumulation and is considered an accelerated state of aging.<sup>371,372</sup> There is increasing evidence suggesting that the increase in and the redistribution of adiposity is the driver of insulin resistance with age, rather than age itself.<sup>11,373,374</sup> Accumulation of visceral white adipose tissue is associated with the development of insulin resistance and cardiometabolic disease,<sup>375</sup> while subcutaneous adiposity is thought to be more metabolically protective.<sup>376</sup> Increased visceral white adipose tissue promotes dyslipidemia and inflammation in the liver, skeletal, and cardiac muscle.<sup>377,378</sup> Moreover, white adipose tissue accumulation leads to a decrease in brown adipose tissue, which also occurs with increasing age (Figure 2; Table).<sup>379,380</sup>

Visceral adiposity also has detrimental effects on the heart, primarily through higher circulating blood volume and increased levels of proinflammatory factors, resulting in elevated stroke volume, cardiac wall stress and injury, and left ventricular hypertrophy.<sup>381</sup> Obesity is, therefore, a major risk factor for developing HF, with each 1-unit increase in body mass index raising the risk of HF by 5% in men and 7% in women,<sup>382</sup> and obesity accounts for ≈20% of atrial fibrillation cases. Accumulation of epicardial fat is of particular importance in the pathogenicity of arrhythmias. Similar body mass indexes may have different CVD risk profiles depending on fat distribution and fitness levels, demonstrating the complex relationship across cardiac, skeletal, and adipose tissues.

## OBESITY AND SKELETAL MUSCLE FUNCTION

Unsurprisingly, energy imbalance worsens skeletal muscle dysfunction in aging. Increased adiposity in older subjects worsened skeletal muscle function independent of age-related decreases in muscle mass, referred to as sarcopenic obesity.<sup>341</sup> Aging and obesity are characterized by chronic inflammation<sup>383,384,385</sup>; therefore elderly individuals with obesity have worsened age-related decreases in muscle strength and mass.<sup>350</sup> These findings are replicated in mouse models, in which both ob/ob and db/db mice have significantly less muscle mass

relative to body mass compared with controls.<sup>386,387,388</sup> Both in vitro and in vivo lipolysis rates are higher in the visceral white adipose leading to increased circulating lipid levels,<sup>389</sup> and these differences are more pronounced in centralized obesity.<sup>390</sup> Increased circulating free fatty acids and adipose-mediated inflammation suppress skeletal muscle fatty acid oxidation and hinder insulin signaling, further impairing systemic glucose homeostasis (Figure 2; Table).<sup>391,392,393</sup>

## OBESITY ALTERS RELEASE OF MYOKINES

Obesity significantly disrupts the production and regulation of myokines, which contributes to metabolic dysfunction. Physical inactivity, a major contributor to obesity, leads to the dysregulation of myokine production, impacting various tissues and metabolic pathways (Figure 2; Table). The changes in myokine levels associated with obesity are complex and often contradictory. For instance, circulating irisin levels are typically lower in individuals with obesity and those with type 2 diabetes,<sup>394,395</sup> but, paradoxically, some studies report elevated irisin under these conditions.<sup>396,215</sup> Exogenous irisin treatment in obese mice has improved glucose tolerance and mitochondrial gene expression,<sup>22</sup> suggesting that elevated irisin levels may, in some cases, reflect compensatory mechanisms or irisin resistance.

Similarly, myostatin, a negative regulator of muscle growth, is elevated in skeletal muscle and serum in obesity, contributing to metabolic dysfunction.<sup>397,398</sup> On the other hand, decreased levels of *Metrnl*, a myokine linked to cardiac health, are associated with impaired cardiac GLUT4 expression and maladaptive cardiac function in hypertrophic and failing hearts.<sup>246,399,400</sup>

In addition, BAIBA, a myokine involved in metabolism, enhances fatty acid oxidation and reduces fat synthesis, showing potential for improving obesity outcomes in mouse models.<sup>401</sup> These findings highlight the complex and context-dependent roles of myokines in regulating metabolism, illustrating their significant impact on the pathophysiology of obesity and related metabolic disorders.

## BARRIERS TO EXERCISE ADHERENCE

Exercise is a cornerstone of health promotion and disease prevention, but, despite its well-documented benefits, adherence to regular physical activity remains a persistent challenge. Barriers to exercise are multifaceted, ranging from personal and psychological to social and systemic influences, highlighting the intricate relationship between these obstacles and overall health.

Exercise is a widely recommended therapeutic intervention to improve health at all stages of life.<sup>402</sup> Despite

extensive efforts to promote this message,<sup>403,404</sup> sedentary lifestyles remain a concern<sup>405,406,407</sup> and contribute to poor health outcomes.<sup>14,408</sup> Poor adherence to an exercise regimen is often due to a variety of barriers, encompassing personal, social, and systemic factors. Common obstacles include lack of perceived time, especially among patients with aging-related conditions,<sup>409,410,411</sup> and limited social support, such as disinterest from friends and family.<sup>409,412</sup> Motivation, energy, and access to resources are often hindered by financial or logistical challenges.<sup>406</sup> Feelings of discomfort, embarrassment, or intimidation in gym settings, particularly among individuals with obesity, further impede participation.<sup>410,413,414</sup> Health issues including chronic diseases, pain, and fatigue, as well as treatments for cancer or autoimmune disease, also reduce adherence.<sup>415,416,417,418</sup> Mental health challenges, including depression, are linked to higher dropout rates, especially in older adults.<sup>419,420,421,422</sup> Socioeconomic constraints, such as inflexible work schedules and cultural norms, also impact exercise adherence.<sup>420,421 423, 424</sup> Collectively, these barriers underscore the complex interplay of factors influencing exercise behaviors. Therefore, addressing factors that contribute to poor exercise adherence is also an essential part of promoting overall global health.

Emerging therapeutic aids may provide additional support for individuals facing exercise adherence challenges. GLP1 (glucagon-like peptide-1) is an exerkine secreted from the gut.<sup>425</sup> In humans, higher cardiorespiratory fitness is associated with lower fasting levels of GLP1 but a higher induction in response to glucose intake.<sup>426</sup> Prescription of GLP1RA (GLP1 receptor agonist) has increased dramatically in recent years due to their effective induction of weight loss. One of the side effects of GLP1RAs is loss of skeletal muscle mass,<sup>427</sup> and exercise could, therefore, be an effective co-adjunct therapy for obesity by counteracting muscle loss and further stimulating metabolic health.

## SUMMARY AND FUTURE PERSPECTIVES

The significant evidence for the health benefits of exercise highlights its critical role in preventing and managing chronic conditions, enhancing cardiovascular and metabolic health, and improving overall quality of life, particularly as populations age. With a deeper understanding of the intricate mechanisms governing skeletal muscle adaptations and the role of exercise-induced signaling molecules, we have an opportunity to develop targeted therapeutic strategies that harness these insights to combat CVD and obesity and promote healthy aging. Future research will be essential in translating these findings into precision exercise interventions that further improve health outcomes across diverse populations.

## ARTICLE INFORMATION

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