

Insights into Sarcopenic Obesity Mechanisms: A Focus on Bidirectional Adipose-Muscle Interactions

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Abstract: Sarcopenic obesity (SO) is a major health issue in aging societies, and its prevalence continues to rise with the global aging process. Compared with sarcopenia or obesity alone, SO is associated with aggravated metabolic disorders and a significantly elevated risk of adverse outcomes. Traditional studies have mostly investigated sarcopenia and obesity as independent pathological conditions; however, recent evidence has revealed a close bidirectional regulatory network between adipose tissue and skeletal muscle, and the imbalance in their crosstalk constitutes the core pathological basis for the occurrence and progression of SO. From the perspective of adipose-skeletal muscle crosstalk, this paper systematically reviews the conceptual framework and bidirectional signaling network in this field, aiming to break through traditional research paradigms and provide an integrated theoretical framework for optimizing diagnostic criteria, identifying biomarkers, and developing precise intervention strategies.

Keywords: sarcopenic obesity, adipo-muscular axis, adipokines, myokines

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1. Introduction

The global population is aging at an unprecedented rate. Population aging has become a major global social transformation in the 21st century, profoundly affecting the economic structures, social security systems and public health policies of countries around the world^[1]. The core characteristic of this phenomenon is the continuous increase in the proportion of the elderly population (generally defined as those aged 60 or 65 years and above) in the total population, accompanied by declining fertility rates and prolonged life expectancy^[2]. Countries around the globe, from China to Greece and from Japan to Poland, are undergoing this structural shift and facing multiple challenges and opportunities brought about by it^[3-6]. Taking Asia as an example, China, the world's most populous country, is experiencing an unprecedented process of population aging. From 2015 to 2050, China has undergone remarkable demographic structural changes, including increased life expectancy and declining fertility rates, which have directly led to population aging^[2]. China's population aged 60 and over is projected to account for 40% of the total population by the middle of this century, marking its entry into a "super-aged society"^[7]. Similarly, 27.4% of Japan's total population is aged 65 and above, and its overall population has been declining continuously since 2006^[6].

With the accelerating process of global population aging, age-related changes in body composition have become an

important issue in public health. Relevant data indicate that population aging and the rising prevalence of obesity show a synchronized increasing trend^[8]. As a global noncommunicable disease, obesity has reached pandemic proportions, affecting more than 890 million people, with an additional 2.5 billion adults being overweight^[9]. Meanwhile, sarcopenia, an aging syndrome characterized by progressive and widespread loss of skeletal muscle mass and function, has gained increasing clinical significance over the past two decades^[10]. When sarcopenia and obesity coexist in the same individual, the condition is defined as sarcopenic obesity (SO), a concept first proposed by Heber in 1996^[11].

Diagnosis of SO requires the simultaneous fulfillment of criteria for both obesity and sarcopenia. However, diagnosis remains challenging, as traditional body mass index (BMI) cannot accurately distinguish between adipose and muscle tissue. Among older adults, severe muscle depletion and excess fat mass may occur even with a normal or high BMI^[12]. Thus, in 2000, Baumgartner provided the first operational definition of SO, defining the sarcopenic component as appendicular skeletal muscle mass divided by height squared falling more than two standard deviations below the mean of a young healthy reference population^[13]. In the subsequent more than two decades, the definition of SO has undergone continuous evolution and debate. In 2022, the European Society for Clinical Nutrition and Metabolism (ESPEN) and the European Association for the Study of Obesity (EASO) jointly issued the first international consensus on the definition and diagnostic criteria for SO^[14]. The consensus recommends a stepwise “screening–diagnosis–staging” workflow: first, obesity is screened using body mass index (BMI) or waist circumference, and sarcopenia is screened based on risk factors or questionnaires; those with positive screening results undergo further assessment of muscle strength. After confirmed low muscle strength, body composition is evaluated by dual-energy X-ray absorptiometry (DXA) or bioelectrical impedance analysis (BIA), with weight-adjusted skeletal muscle mass and body fat percentage used to define sarcopenia and obesity, respectively. The publication of this consensus marks a critical step toward standardization and comparability in the field of SO research.

In recent years, large-scale population-based meta-analyses have shown that the global prevalence of SO in the elderly population is approximately 7%. However, the reported prevalence varies from 2.1% to 12% due to differences in diagnostic criteria and population characteristics^[15]. Of note, SO is not a simple coexistence of sarcopenia and obesity. A growing body of evidence indicates that the combination of these two conditions exerts synergistic adverse health effects. A meta-analysis including nearly 580,000 participants demonstrated that SO was significantly associated with a 2.06-fold increased risk of cardiovascular and cerebrovascular diseases^[16]. In the field of oncology, SO has also been confirmed to be closely associated with shortened overall survival (HR = 1.52) and an increased risk of postoperative complications (OR = 2.23) in patients with non-metastatic colorectal cancer^[17]. These findings suggest that the health threat posed by SO is far greater than that of sarcopenia or obesity alone. Traditionally, adipose tissue and skeletal muscle have been regarded as functionally relatively independent organs. However, recent studies have revealed that a close bidirectional regulatory network exists between them, forming the so-called “Adipo-Muscular Axis”^[19]. Adipose tissue and skeletal muscle are not only organs for energy storage and expenditure, but also active endocrine organs. The adipokines and myokines they secrete form the chemical language for inter-tissue communication, regulating metabolic homeostasis through autocrine, paracrine, and endocrine pathways^[18, 19]. Dysregulation of this interactive axis constitutes the core pathological basis of SO: dysfunctional adipose tissue in obesity leads to increased secretion of pro-inflammatory adipokines, free fatty acid spillover, and ectopic lipid deposition, which in turn trigger insulin resistance and mitochondrial damage in skeletal muscle. Conversely, reduced muscle mass and diminished muscle strength lower resting energy expenditure, further exacerbating fat accumulation and forming a vicious cycle^[20, 21]. This article aims to systematically review the central role of bidirectional crosstalk between adipose tissue and skeletal muscle in the development and progression of SO, with a focus on the molecular mechanisms mediated by intermuscular adipose tissue, intramuscular adipose tissue, adipokines and myokines, and to discuss the implications of this novel conceptual framework for the identification of SO biomarkers and the development of precise intervention strategies.

2. Anatomical and Functional Basis of Adipose-Skeletal Muscle Crosstalk

Adipose tissue and skeletal muscle are closely adjacent in anatomical structure, and this spatial proximity provides a structural basis for their functional crosstalk. According to anatomical location, adipose tissue is distributed in multiple functionally and metabolically heterogeneous fat depots throughout the body. Among them, those directly associated with skeletal muscle are intermuscular adipose tissue (IMAT) and intramyocellular lipid (IMCL).

IMAT refers to adipose tissue located beneath the muscle fascia and between muscle bundles, constituting a unique anatomical fat depot within the musculoskeletal system^[22]. Unlike subcutaneous and visceral adipose tissue, IMAT is adjacent to the epimysium, with adipocytes accumulating in the connective tissue septa between muscle groups. Although IMAT is separated from muscle fibers by fascia, the adipokines and bioactive molecules it secretes can still act on adjacent muscle tissue through paracrine pathways, regulating the local metabolic microenvironment^[23]. Several studies have suggested that IMAT may negatively regulate insulin sensitivity and promote insulin resistance by secreting free fatty acids (FFA), inflammatory factors (such as PAI1 and MCP1), and other bioactive molecules^[24]. From the perspective of cellular origin, the formation of IMAT is closely associated with the aberrant differentiation of fibro/adipogenic progenitors (FAPs). FAPs are a highly plastic cell population residing in the skeletal muscle interstitium. Under physiological conditions, they interact with muscle satellite cells and immune cells via paracrine signaling, providing a supportive microenvironment for muscle regeneration^[25]. However, in the context of aging, obesity and metabolic diseases, the fate of FAPs is regulated by a complex network of multiple factors including the Wnt signaling pathway and PPAR γ , thereby leading to the abnormal expansion and accumulation of intermuscular adipose tissue^[26].

IMCL is located within muscle fibers and serves as an intramyocellular lipid storage depot. It directly infiltrates the muscle microenvironment and represents the main component of muscular fatty infiltration^[27]. Adipocytes within IMCL are in close proximity to muscle fibers without separation by fascia. This high spatial proximity allows IMCL to directly regulate muscle fiber function through dual pathways: physical compression and paracrine signaling. On the one hand, abnormal accumulation of IMCL can disrupt the normal arrangement of muscle fibers and impair the mechanical efficiency of muscle contraction^[28, 29]. On the other hand, as an active endocrine tissue, IMCL can secrete a variety of pro-inflammatory adipokines (such as TNF- α and IL-6) and chemokines, which act on adjacent muscle fibers through paracrine pathways, inducing local inflammatory responses and insulin resistance^[30, 31]. From the perspective of vascular-muscle crosstalk, abnormal expansion of IMCL can exert mechanical compression on blood vessels running intermuscularly or intramuscularly, increasing vascular resistance and reducing microcirculatory perfusion^[32]. This process directly limits the supply of oxygen and metabolic substrates to myocytes, constituting an important vascular mechanism underlying the decline in muscle function. Similar to IMAT, FAPs are also a major source of IMCL. Under pathological conditions, FAPs shift from a functional state that supports muscle regeneration to an adipogenic phenotype, resulting in the accumulation of adipocytes within muscle fibers. Studies by Biltz et al. have confirmed that intramuscular adipose tissue infiltration impairs skeletal muscle contractile function, reduces muscle strength, and exacerbates muscle fibrosis^[33].

Traditionally, adipose tissue has been regarded as a passive energy storage organ, and skeletal muscle as an active contractile organ. In recent years, a growing body of research has confirmed that adipose tissue and skeletal muscle are not only central organs for energy storage and expenditure, but also active endocrine organs. Through the secretion of various bioactive factors, they engage in extensive and complex crosstalk, jointly regulating systemic energy metabolism, insulin sensitivity, inflammatory status, and tissue remodeling^[34-36]. On this basis, researchers have proposed the conceptual framework of the “Adipo-Muscular Axis” to characterize the bidirectional regulatory network between the two tissues^[37]. The core connotations of this axis include: (1) Adipose tissue regulates skeletal muscle protein homeostasis, mitochondrial function, and insulin sensitivity via adipokines, extracellular vesicles, and lipid metabolites; (2) Skeletal muscle reciprocally modulates adipocyte differentiation, thermogenic function, and lipid metabolism through myokines; (3) The integrated effect of their crosstalk maintains systemic energy metabolic homeostasis.

Under physiological conditions, the adipo-muscular axis maintains a dynamic balance: adipose tissue secretes appropriate regulatory signals to ensure energy supply and functional maintenance of skeletal muscle; skeletal muscle

reciprocally modulates the metabolic adaptation of adipose tissue through contractile activity and myokine secretion. However, in the context of obesity, aging and metabolic diseases, this balance is disrupted, leading to a vicious pathological cycle. In obesity, adipose tissue dysfunction results in increased secretion of pro-inflammatory adipokines, free fatty acid spillover, and ectopic lipid deposition^[38, 39]. Accumulation of lipid metabolic intermediates (such as ceramides and diacylglycerols) in myocytes disrupts insulin signaling pathways and induces mitochondrial dysfunction, leading to skeletal muscle insulin resistance and enhanced catabolism^[40, 41]. Meanwhile, the loss of skeletal muscle mass and decline in muscle strength reduce resting energy expenditure, further exacerbating fat accumulation^[42]. The vicious cycle in which fat accumulation and muscle function decline reinforce each other as cause and effect constitutes the core pathological basis of SO.

3. Mechanisms of Adipose Tissue Action on Skeletal Muscle

Adipose tissue is composed of multiple cell types, including adipocytes, fibroblasts, endothelial cells, immune cells, and so on. These cells collectively secrete a diverse array of bioactive molecules that constitute the adipose secretome^[43]. Adipokines represent the most extensively studied group, including leptin, adiponectin, resistin and others. They can act on distant organs through the endocrine pathway, and also regulate the function of adjacent tissues via paracrine signaling^[44]. In obesity, the secretory profile of adipose tissue undergoes a significant shift from an anti-inflammatory to a pro-inflammatory phenotype, with increased secretion of pro-inflammatory adipokines and decreased levels of protective adipokines.

According to their functional characteristics, adipokines can be categorized into two major groups: proinflammatory adipokines and protective adipokines. The imbalanced expression of these two groups in obesity is a key feature of adipomuscular axis dysfunction: (1) Proinflammatory adipokines: TNF α and IL6 are the most extensively studied proinflammatory adipokines. In obesity, adipose tissue exhibits increased macrophage infiltration, resulting in markedly upregulated secretion of TNF α and IL6^[45]. TNF- α can induce the expression of the E3 ubiquitin ligases atrogin-1 and MuRF1 in skeletal muscle cells by activating the NF- κ B and p38 MAPK signaling pathways, thereby promoting protein degradation in muscle fibers^[46]. Meanwhile, TNF- α can also interfere with insulin signal transduction by inhibiting tyrosine phosphorylation of insulin receptor substrate1 (IRS1), leading to insulin resistance in skeletal muscle^[47]. IL6 inhibits the critical muscle anabolic pathway of IGF1/PI3K/Akt/mTOR by activating the SOCS3 signaling pathway^[48, 49]. IL-6 also impairs glucose uptake in skeletal muscle, leading to metabolic dysfunction^[50]. Existing studies have shown that IL-6 levels are significantly elevated in patients with SO and are positively correlated with disease severity^[51]. In addition, leptin is also one of the most representative adipokines secreted by adipose tissue. Under physiological conditions, leptin exerts its core effects of regulating appetite, increasing energy expenditure, and maintaining energy homeostasis by binding to its hypothalamic receptors and activating the JAK2/STAT3 signaling pathway^[52]. However, in obesity, despite markedly elevated circulating leptin levels (hyperleptinemia), leptin resistance often occurs^[53]. In the resistant state, hyperleptinemia not only fails to effectively suppress appetite, but also promotes the production of pro-inflammatory cytokines such as TNF- α and IL-6 by activating immune cells including T cells and macrophages, thereby inducing systemic low-grade inflammation^[54]. Studies have confirmed that serum leptin levels are significantly associated with SO. Even after adjustment for multiple confounding factors, leptin levels remain independently positively correlated with BMI and independently negatively correlated with skeletal muscle index (SMI)^[55]. (2) Protective adipokines: Unlike most adipokines, adiponectin is an adipokine with anti-inflammatory, insulin-sensitizing, and metabolic regulatory effects, whose levels are conversely decreased in obese individuals^[56]. Adiponectin promotes fatty acid oxidation, inhibits hepatic gluconeogenesis, and improves insulin sensitivity in skeletal muscle by activating the AMPK and PPAR- α pathways^[57, 58]. Adiponectin also inhibits the release of pro-inflammatory cytokines such as TNF- α and IL-6, thereby alleviating chronic low-grade inflammation^[59]. A cross-sectional study showed that low adiponectin levels are associated with poorer muscle function and physical performance in patients with sarcopenia^[60]. Extracellular vesicles (EVs), as novel mediators of

intercellular communication, have attracted extensive attention in the field of adipose-skeletal muscle crosstalk research in recent years. EVs are nano to micrometersized particles with a lipid bilayer membrane structure, which can carry bioactive molecules such as proteins, microRNAs, and lipids to transmit signals between tissues and regulate metabolic homeostasis^[61]. EVs derived from adipose tissue (especially white adipose tissue, WAT), termed AdEVs, can reach skeletal muscle through the circulatory system and affect its metabolism and function^[62]. Recent studies have shown that miR-146a-5p carried by adipose tissue-derived small extracellular vesicles (sEVs) can target Fbx32 (Atrogin-1) to inhibit excessive activation of mitophagy, reduce apoptosis and reactive oxygen species (ROS) production, and enhance ATP generation, thereby modulating the progression of muscle aging and muscle atrophy^[63]. Adipose-derived sEVs also carry various other miRNA molecules, such as miR-27a, which can participate in the regulation of skeletal muscle lipid metabolism by targeting signaling pathways including PPAR γ ^[64]. An animal study found that EVs derived from obese mouse models significantly upregulated the secretion of IL6 (a proinflammatory factor) and inhibited the secretion of IL4 (an antiinflammatory factor) in a coculture system, suggesting that obesity-associated EVs may promote muscle degradation through a proinflammatory microenvironment^[65]. These findings suggest that adipose tissue constitutes a novel molecular pathway regulating skeletal muscle function through sEV-mediated miRNA delivery, providing a new perspective for understanding the mechanism of adipose-skeletal muscle crosstalk.

4. Retrograde regulation of adipose tissue by skeletal muscle

Skeletal muscle also possesses active endocrine function and can synthesize and secrete a variety of myokines. Myokines are cytokines or polypeptides secreted by skeletal muscle, which regulate multi-organ function through autocrine, paracrine or endocrine modes^[66]. Released after exercise, they mediate the beneficial effects of exercise on metabolism, immunity and energy homeostasis^[67]. Several myokines have been identified to date, including irisin, brain-derived neurotrophic factor (BDNF), myostatin (MSTN), and meteorin^[68]. These molecules play central roles in regulating the function of adipose tissue. Irisin is produced through the exercise-induced PGC-1 α /FNDC5 pathway, promotes the conversion of white adipocytes into beige fat by activating UCP1 expression in these cells, enhances thermogenesis and energy expenditure, and improves glycolipid metabolism in obesity and diabetes^[69, 70]. Irisin can significantly inhibit adipogenic differentiation of adipose-derived mesenchymal stromal cells (ASCs), reduce their antioxidant capacity, and simultaneously increase ATP production and the generation of reactive oxygen species (such as superoxide anion), suggesting its dual regulatory role in adipose tissue homeostasis^[71]. Studies have confirmed that irisin levels are negatively correlated with BMI and body fat percentage, and may serve as a predictor of adipose dysfunction^[72]. Myostatin (also known as GDF8), a member of the transforming growth factor β superfamily primarily secreted by skeletal muscle, is traditionally regarded as a negative regulator of muscle mass^[73]. However, recent studies have revealed that it also plays a similarly complex and critical role in the metabolism and functional remodeling of adipose tissue, as well as in the regulation of systemic energy homeostasis^[74]. Studies have found that myostatin is not secreted exclusively by muscle; brown and white adipose tissues themselves can also express and secrete a certain amount of myostatin^[75]. Myostatin derived from brown adipose tissue exhibits autocrine and paracrine functions, thereby regulating the thermogenic function of brown adipose tissue itself^[76]. For example, specific knockout of the myostatin gene in brown adipose tissue results in increased body weight, hepatic steatosis, and is accompanied by mitochondrial dysfunction and inflammatory characteristics in brown adipose tissue in mice fed a high-fat diet^[76].

In addition to the direct effects of myokines, the overall metabolic status of skeletal muscle affects adipose tissue function through various indirect pathways. As the major consumer of systemic glucose and fatty acids, the metabolic activity of skeletal muscle directly determines the flow of energy substrates. Under insulin-sensitive conditions, skeletal muscle efficiently takes up and oxidizes glucose and fatty acids, reducing lipid overflow into adipose tissue; in contrast, under insulin-resistant conditions, the metabolic load of skeletal muscle decreases, redirecting more energy substrates to adipose tissue for storage and exacerbating obesity^[77]. Furthermore, the protein turnover status of skeletal muscle affects

adipose tissue function through amino acid metabolism. Increased muscle protein synthesis enhances the utilization of branched-chain amino acids (BCAAs) and reduces circulating BCAA levels, whereas enhanced muscle breakdown leads to BCAA overflow, which activates mTORC1 signaling in adipocytes and induces insulin resistance in adipose tissue^[78].

5. Conclusion

From the perspective of adipose-skeletal muscle crosstalk, this article systematically reviews the bidirectional signaling network underlying SO. Adipose tissue acts on skeletal muscle through multiple pathways, including pro-inflammatory adipokines, extracellular vesicles, and lipid metabolites, disrupting insulin signaling, inducing mitochondrial dysfunction, and promoting protein degradation; in turn, skeletal muscle reciprocally regulates the differentiation and thermogenic function of adipose tissue via myokines. Imbalance in this bidirectional network leads to fat accumulation, impaired muscle function, and reduced energy expenditure, forming the core pathological basis of SO. Future research should further explore the regulatory mechanisms of extracellular vesicles as novel mediators, the functional significance of lipid droplet heterogeneity, multi-omics-based molecular classification strategies, and develop precise interventions targeting the adipose-skeletal muscle axis. The paradigm shift from “dual-organ crosstalk” to a “multi-system network” will open new avenues for the early diagnosis and treatment of SO.

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