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Microfluidic systems for studying dynamic function of adipocytes and adipose tissue

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Abstract

Recent breakthroughs in organ-on-a-chip and related technologies have highlighted the extraordinary potential for microfluidics to not only make lasting impacts in the understanding of biological systems but also to create new and important in vitro culture platforms. Adipose tissue (fat), in particular, is one that should be amenable to microfluidic mimics of its microenvironment. While the tissue was traditionally considered important only for energy storage, it is now understood to be an integral part of the endocrine system that secretes hormones and responds to various stimuli. As such, adipocyte function is central to the understanding of pathological conditions such as obesity, diabetes, and metabolic syndrome. Despite the importance of the tissue, only recently have significant strides been made in studying dynamic function of adipocytes or adipose tissues on microfluidic devices. In this critical review, we highlight new developments in the special class of microfluidic systems aimed at culture and interrogation of adipose tissue, a subfield of microfluidics that we contend is only in its infancy. We close by reflecting on these studies as we forecast a promising future, where microfluidic technologies should be capable of mimicking the adipose tissue microenvironment and provide novel insights into its physiological roles in the normal and diseased states.

Keywords

Microfluidics microfabrication; Adipocyte; Adipokine; Batokine; Hormone; Bioanalytical methods

Introduction

The global epidemic of obesity has brought increasing attention to research for understanding the biology of adipocytes (fat cells). Adipose tissue or fat tissue, which accounts for 5% to 50% of human body weight, was traditionally thought to be a passive reservoir for long-term energy storage, but now it is known to be a complex, essential, and highly active metabolic and endocrine organ with meaningful roles in the integration of systemic metabolism [1]. There are two major types of mammalian adipose tissues, white adipose tissue (WAT) and brown adipose tissue (BAT) [2], although intermediate (e.g., "brite" or "beige" cells) and locale-specific subtypes (visceral, omental, etc..) do exist.

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WAT is the main type of adipose tissue in the adult human and is mainly found in subcutaneous regions and visceral depots. WAT consists of adipocytes (the main cellular component of WAT that are crucial for both energy storage and endocrine activity), precursor cells, fibroblasts, vascular cells, macrophages, smooth muscle, and immune cells. WAT is a critical regulator of system energy homeostasis and acts both for energy storage and as an endocrine organ. Under the appropriate hormone levels (e.g., high insulin), extra energy obtained from food or from the bloodstream, including glucose and fatty acids, is absorbed by adipocytes and transformed into triacylglycerols during lipogenesis for longterm storage. Upon energy deficiency or during other hormonal signaling events (e.g., low insulin, high glucagon), triacylglycerols are hydrolyzed (lipolysis), and the non-esterified fatty acid (NEFA) and glycerol components are released into circulation for delivery to other organs. The processes of glucose and fatty acid uptake, lipogenesis, and lipolysis are precisely controlled by a complex regulatory network, a topic which is reviewed elsewhere [3, 4].

WAT also secretes numerous hormones that regulate the metabolic functions of nearly all other organs; these hormones are referred to as "adipokines." A detailed list of key adipokines and their known sources and functions are summarized in Table 1. More importantly, under the condition of obesity, WAT not only accumulates and expands throughout the body but also recruits immune cells that infiltrate into adipose tissue and change the adipokine secretion profile. Although this process is not fully understood, it is related to systemic inflammation and insulin resistance [20].

Distinctly differing from WAT, the BAT derives its signature brown color from the high number of mitochondria, and the tissue mainly participates in thermogenesis. BAT helps maintain normal body temperature in newborn infants and hibernating mammals, and while its percentage of total body mass decreases with age, it is still present in the normal adult human [21]. This tissue is also known to have systemic effects by secreting regulatory molecules, referred to as batokines [22], and the specific batokine profile is distinct from adipokines. Batokines with autocrine, paracrine, and endocrine effects are listed in Table 2.

Although our understanding of adipose tissue biology has greatly improved over the last few decades, the level of established detail generally lags behind that of other endocrine systems. Consequently, researchers continue to actively experiment on the tissue and its constituent cells in attempts to further elucidate its functions and mechanisms, to better understand the pathogenesis of obesity-related disorders, and to develop therapeutic strategies for these disorders. This heightened interest has catalyzed a demand for higher performance bioanalytical methodology, and several groups—including our own—have recently shown microfluidics to offer unique opportunities for studies on adipose tissue [39–52]. Microfluidic analysis systems offer a host of well-established benefits, including decreases in experiment cost and reagent volume, increased temporal and spatial resolution in cellular function assays, and in vivo mimics of vasculature and micro-environments within channels and chambers on the devices [53]. However, when applying microfluidics to studying adipose tissue, unique challenges such as cell buoyancy give rise to specialized countermeasures that arguably position such devices into their own subcategory. In this review, we highlight recent developments in the special class of bioanalytical microfluidic

devices designed to study the dynamics of adipocytes and adipose tissue toward a better understanding of the biology related to obesity, diabetes, and metabolic disorders.

Microfluidic platforms to study adipocytes

In-vitro cell models integrated onto microfluidic devices

Historically, most adipocyte-related studies—including adipocyte differentiation, lipogenesis, adipokine secretion, and regulation—have been carried out with immortalized cell lines, such as mouse 3T3-L1 [54], 3T3-F442A [55], or human Simpson-Golabi-Behmel syndrome (SGBS) cells [56]. Differentiation of these cells into adipocytes can be achieved in vitro by treatment with adipogenic stimuli, including cAMP agonists, insulin, and glucocorticoids [57]. These cell lines share many similarities with primary adipocytes such as fat storage in lipid droplets, insulin sensitivity, expression of adipocyte-specific genes, and adipokine secretion [2]. However, there are some important differences. The primary white adipocyte from a host mammal contains a single large lipid droplet, but the cell lines exhibit clustering of many smaller droplets. Also, the cell line's adipokine secretome differs from that of primary adipocytes; for example, leptin expression is at much lower levels in 3T3-L1 and 3T3-F442A cells compared with primary tissue [58].

Recently, the development of tissue engineering technology has allowed assembly of functional tissues with multiple cells and IGF13D extracellular matrix materials to recapitulate many of the key features of both normal and pathological human organs in vitro. Researchers have successfully constructed those organ-on-a-chip models for the study of liver, kidney, intestine, lung, heart, smooth and striated muscle, fat, bone, marrow, cornea, skin, blood vessels, nerve, and blood-brain barrier over the past decade [59]. Various types of engineered adipose tissues were fabricated by incorporation of adipose cell lines into extracellular matrix materials including collagen [41, 60], silk fibroin [61], sponge-like porous polyurethane [62], decellularized extracellular matrices [35], and hyaluronic acidbased hydrogels [36]. These novel in vitro models showed better simulation of functions of in vivo adipose tissue than cell lines cultured with traditional methods. Engineered adipose tissues were also used as a simplified model to address several challenges in organ-on-a-chip engineering. For example, the importance of the "right size" of the tissue for maintaining in vivo basal metabolic rates was studied by measurement of insulin-induced glucose uptake by collagen confined adipose spheroids with various sizes [41]. Transport processes by diffusion through porous membranes within multilayer microfluidic devices, providing shear stress protection to cells or tissues cultured on-chip, were also visualized by fluorescently labeled fatty acid analog uptake by on-chip differentiated 3T3-L1 adipocytes [40]. The Meier group used a microfluidic large-scale integration (mLSI) culture platform with integrated assays to study important factors in stem cell differentiation into adipoocytes [63]. However, since this review is focused on micro-analytical systems that study the dynamics of adipose tissue metabolism, hormone secretion, and other functions, approaches using such bottom-up adipocyte-on-a-chip engineering, which has been reviewed elsewhere [64], is only covered where appropriate.

Integrating primary tissue or pre-differentiated adipocytes onto microfluidic systems

While the cell lines and engineered adipose tissue remain as valuable in vitro models, much of the adipocyte's in vivo function can only be recapitulated through primary cultures. Primary adipose tissues can be isolated from lab animals and cultured according to well established protocols [42]. The adipose tissue pad or explant can be taken directly from a donor and directly integrated onto microfluidic devices, or it is often beneficial to digest the matrix tissue within the explant and isolate constituent primary adipocytes, to avoid the interference from other cell components.

Owing to the inherent positive buoyancy of the lipid droplets already present in primary tissue or pre-differentiated adipocytes, dispersed cells or tissue explants must be properly trapped or anchored within microfluidic devices. This approach prevents floating of the cells in aqueous media, allows controlled fluidic access to and from the tissue, and avoids possible clogging of microfluidic channels. This buoyancy issue has been addressed in several ways by our research group recently, including trapping cells with metallic mesh [39], 3D-printed accessories [44] (Fig. 1-C), and collagen matrix casting [45]. The issue has been addressed by others using a culture chamber with a micro-post array [52], integrating cell-laden coverslips onto multilayer devices [46, 47], or by other methods as shown in Table 3. As shown in the table, within the past several years, researchers have devised a number of trapping methods that can be chosen by those interested in using microfluidics to study adipose tissue dynamics.

Dynamic adipose tissue function evaluated through microfluidics

Insulin-induced glucose uptake

Blood glucose levels are precisely regulated by the hormones insulin and glucagon within a narrow, normal range via strong endocrine signaling throughout the body. During feeding, insulin (secreted by pancreatic islets) increases glucose uptake into adipose and muscle tissues through glucose transporter type 4 (GLUT4) [65], a high affinity glucose transporter mostly expressed in these tissues. In the absence of insulin, only 5% of the GLUT4 is found at the cell surfaces. Insulin recruits GLUT4 to translocate to the plasma membrane from specialized GLUT4 storage vesicles, leading to a concomitant increase in cellular glucose uptake [66]. Although adipose tissue only accounts for about 10% of the insulin-stimulated glucose uptake (the other 90% occur in skeletal muscle) [67], this process is very important for energy homeostasis, as the secreted adipokines from WAT also regulate whole-body metabolism as stated earlier. Therefore, understanding the dynamics of regulated glucose transport in adipocytes using microfluidic methods should provide better opportunities for elucidating the physiological and pathophysiological mechanism of energy homeostasis.

The Elvassore group reported the first microfluidic human adipose tissue glucose uptake study [52]. Adipose tissue was trapped on an automated microfluidic injection system, which allows time resolved insulin stimulation. By comparing tissues from Type 2 diabetic and non-diabetic patients, insulin-resistance effects were observed in diabetic tissues when biopsies were treated with an insulin and glucose step, mimicking the postprandial phase. The healthy tissue showed a significant increase in glucose uptake upon insulin stimulation,

whereas the diabetic tissue had no evidence of a difference in glucose uptake rate. However, the authors applied only one treatment to each of the precious tissue samples, even though the device was said to allow 1 wk of on-chip tissue culturing. Furthermore, the glucose concentration in the outflow was only measured every 10 min even though their system should be capable of ~2 min interrogation.

In a particularly valuable in vitro model for metabolic study, 3T3-L1 adipocyte spheroids were used by the Takayama group to study the scaling effects in a human-on-a-chip system [41]. They highlight the fact that allometry—the relationship between body size and shape, including anatomy and physiology—is a significant factor in organ-on-a-chip system design that should be considered when estimating the appropriate size of the "chip organism" for maintaining a similar basal metabolism rate (BMR) as in vivo. The insulin-induced glucose uptake rates by intact 3T3-L1 spheroids or mechanically dispersed spheroids within collagen were compared. This work demonstrated a significant glucose uptake difference between intact and mechanically dispersed spheroids, establishing that tissue architecture can significantly affect scaling relationships in microfabricated devices.

Fatty acid uptake

The energy reserves in adipose tissues are stored as triacylglycerol (TAG) molecules, which are mainly obtained from two routes: de novo lipogenesis from non-lipid precursors such as glucose, or uptake of free fatty acid and TAG from circulating blood. As the solubilities of free fatty acids and TAG in aqueous solution are extremely low [68], fatty acids or TAG molecules in blood are carried by lipoproteins including chylomicrons (CM), very lowdensity lipoproteins (VLDL), and serum albumin. Lipoprotein lipase (LPL), which is a glycoprotein secreted by adipocytes and translocated to the lumen of endothelial cells, digests TAG bound to lipoproteins thereby releasing free fatty acid for uptake by adipocytes [69]. The transport of non-esterified fatty acids (NEFAs) across the cell membranes is promoted by CD36/SR-B2, which is currently known as the predominant membrane protein facilitating fatty acid transport in adipocytes, enterocytes, cardiac myocytes, and skeletal myocytes [3]. Insulin-induced CD36/SR-B2 translocation from endosomal compartments to the cell membrane results in upregulated fatty acid uptake [3], which is a mechanism very similar to the GLUT4 regulation discussed above.

In our recent study, fatty acid uptake and release was monitored by fluorescence microscopy imaging of adipose tissue explants confined within our novel 16-channel microfluidic multiplexer (μMUX) [39]. Adipose tissue explants were exposed to temporal mimics of post-prandial insulin and glucose levels while simultaneously switching between fluorescently labeled and unlabeled free fatty acids. This unique experiment, enabled by the microfluidic platform, permitted fluorescence imaging of fatty acid uptake dynamics in real time (Fig. 1D–H). Surprisingly, both the initial rates of fatty acid uptake and the release rates were observed to follow the pattern of glucose and insulin. Moreover, treatment with a CD36/SR-B2 inhibitor [70] suppressed both the fatty acid uptake and release [43]. These results suggest the current understanding of fatty acid uptake and regulation is incomplete. It is likely that CD36/SR-B2 is not only involved in fatty acid uptake but plays a key role in fatty acid release. Based on our results [39, 43], this fatty acid exchange equilibrium

between uptake and release is likely regulated by insulin induced CD36/SR-B2 translocation as well as by the extracellular fatty acid concentration. This hypothesis also agrees with our observation of fatty acid transportation between adipocytes, where adipocytes along the edges of the explants absorbed fatty acids more rapidly than inner cells. Over time, fatty acid could be seen permeating into centrally positioned adipocytes [43]. Further experiments to test this hypothesis are ongoing in our group, where the microfluidic platforms shown in Fig. 1C–H are key components that provide unique experimental capabilities to ask these questions.

Secretion of non-esterified fatty acids and glycerol

In the starving or fasting states, TAGs are hydrolyzed into NEFAs and glycerol by the lipase process, and the products are secreted into blood circulation by adipose tissues as an energy source for other tissues. The hydrolysis of stored TAG involves several lipases, including hormone-sensitive lipase (HSL), monoacylglycerol lipase (MGL), and adipose triglyceride lipase (ATGL) [71]. TAG is hydrolyzed by ATGL to diacylglycerol (DAG) and one molecule of fatty acid. HSL then converts DAG to a second fatty acid and monoacylglycerol (MAG). In the last step, MGL hydrolyzes MAG to glycerol and the third fatty acid molecule. The NEFAs and glycerol are not only used as nutrients for the rest of the body, but they also have important signaling effects on energy homeostasis. For example, circulating NEFAs have been shown to reduce glucose uptake by adipocytes and muscle and increase hepatic gluconeogenesis [72], and chronic exposure to elevated level of NEFAs was shown to decrease beta cell insulin secretion [73].

Since their 2009 publication of the seminal work in the area [46], the Kennedy group has reported several microfluidic studies where NEFA and glycerol secretion was quantified from 3T3-L1 adipocytes. The 3T3-L1 adipocytes were pre-differentiated onto cover slips and integrated into a reversibly sealed multilayer microfluidics system. The constant flow containing the secretomes was mixed with NEFA or glycerol enzyme probes, and the concentrations of NEFAs and/or glycerol were detected by fluorescence imaging on-chip [46–48, 50]. Upon treatment with a beta-adrenergic agonist, up to 6-fold, sustained increases in NEFA and glycerol secretions were observed. The latest version of their microfluidic device (Fig. 1A, B) has combined solid phase extraction and mass spectrometry, providing unique dynamic information about complex NEFA secretion profiles from adipocytes [49]. Use of such integrated microsystems—combining adipocyte culture, sample preparation, microfluidic sampling, and mass spectrometry—has the potential to make significant impacts to our understanding of adipose tissue biology.

Our group also has studied glycerol secretion from primary adipose tissue explants [44]. In order to counteract the adipose pad buoyancy, in this work we used custom 3D-printed accessories to trap the tissues into on-chip reservoirs. After 30 min of high-insulin/highglucose treatments, the explants were switched to low-glucose/low-insulin solution, and increased glycerol secretion rates were observed.

Adipokine or batokine secretion

As stated earlier, adipose tissue plays a significant role in the endocrine system for maintaining whole body energy homeostasis. However, to the best of our knowledge, our group has conducted the only adipokine secretion study using microfluidics [45]. In this design, adipocytes from dispersed primary mouse tissue were cultured in collagen within a customized culture reservoir using a raised island to interface to the microchannel. Secreted adiponectin, which is a large multimeric protein hormone with the main understood function of improving insulin sensitivity, was quantified after sampling from this device. Insulin and niacin treatments induced increased adiponectin secretion by 2.6- and 4.4-fold, respectively.

Considering the importance of adipokine and batokine secretion in physiology, we assert that microfluidics should be an ideal tool for future studies on the dynamics of adiposederived hormone secretion and its downstream systemic effects. As such, the lists of adipokines in Table 1 and batokines in Table 2 are provided also as a reference for those interested in applying microfluidic systems to important problems in the area of adipose tissue biology.

Thermogenesis

Heat generation is one of the necessities for endotherms to maintain normal body temperature and routine metabolism. BAT is known as the main site of non-shivering thermogenesis, which enables animals to adapt to a cold environment. Uncoupling protein 1 (UCP1) is a transmembrane protein only expressed in brown adipocytes. UCP1 is located on the inner membrane of mitochondria, where it increases the inner membrane proton permeability, decreases the proton gradient generated in oxidative phosphorylation, uncouples the respiratory chain, and yields a higher oxidation rate for heat generation [22]. UCP1-mediated proton leakage is regulated by the sympathetic nervous system through βadrenergic receptors, and binding of catecholamines (such as norepinephrine) to βadrenergic receptors in BAT trigger thermogenesis. UCP1 is also activated by fatty acids and inhibited by purine nucleotide di- and triphosphates (e.g., ATP, ADP, GTP, and GDP).

To investigate the heat generation property of BAT, Inomata et al. [51] fabricated a microfluidic system with silicon resonator-based thermal sensors, facilitating thermal detection of single cells. Heat generated from single brown adipocyte cells was measured by resonant frequency changes in the resonators. With their devices, two types of heat emissions from single brown fat cells were detected. Continuous heat generation during norepinephrine stimulation was quantified, and pulsed heat generation was quantified without any stimulation. Cells treated with sodium azide, a respiratory inhibitor, did not show any thermal response. This study represents the only report, to our knowledge, that has assayed dynamic function of BAT using microfluidics.

Conclusions and future outlook

Herein, we have reviewed some of the recent microfluidic systems applied to the study of adipocytes, or fat cells. As summarized in Table 3, these studies include adipocyte nutrient uptake, fatty acid and glycerol secretion, hormone secretion, and thermogenesis. From these

reports, we have observed that microfluidic platforms can provide high precision analytical tools to promote improved understanding of adipocyte biology. Several new microfluidicsenabled findings—particularly the time resolved NEFA secretion profile, adiponectin secretion, and the first real time observation of insulin regulated fatty acid exchange—have provided new information on the dynamics of adipose tissue function. Future work in this new sub-field of microfluidics has the potential to inform better treatments for obesity, diabetes, and metabolic disorders.

Compared to the relatively effective applications of microfluidic systems to other endocrine tissues or cell types in recent years, such as islets of Langerhans [74, 75], far fewer studies have been conducted on adipose tissue using microfluidics thus far—especially on brown adipose tissue. We surmise that there are several reasons for this deficiency in the literature. First, the importance of adipocytes in energy homeostasis, and particularly the endocrine functions of adipose tissue, has been less explored, thereby reducing exposure of the concepts to the microfluidics community. In fact, a chief purpose of providing this critical review is to address this issue. Secondly, the buoyancy of adipocytes introduces difficulties to incorporate these cells onto microfluidic devices. To prevent cell loss and channel clogging, device engineers must use creative ways to trap or anchor the cells on chip, such as the 3D-printing technology used by our group for cell-driven macro-to-micro interfacing. Thirdly, the fundamental understanding of adipocytes is still relatively underdeveloped. For example, several new hormones were identified as recently as 2016 [76], namely a new adipokine (asprosin) and two new batokines (Slit2-C and PM20D1). These latest findings, along with the more complete lists of adipose-derived hormones (Table 1 and Table 2) represent great opportunities for bioengineers and bioanalytical chemists to make contributions to the adipose biology field.

Further innovations in microfluidics should continue to provide a unique toolset that allows biologists to advance their research in ways that were not previously possible. Conversely, bioengineers and bioanalytical scientists are also seeking unique problems in medicine and biology to validate and improve their microfluidic technologies. As stated in this review, the relatively understudied adipose tissue could be an important target tissue for the microfluidics community. Concurrently, further improvements to conform microfluidics into adipose tissue biology should provide unique opportunities toward better preventions and treatments, meeting urgent needs in the global epidemics of obesity, diabetes, and metabolic disorders.

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Fig. 1.

Recent examples of microfluidic systems for studying dynamic function of adipocytes and adipose tissue. (**A**) Schematic of microfluidic device for 3T3-L1 adipocyte stimulation and sampling, coupled to solid phase extraction and mass spectrometry (SPE-MS), for monitoring non-esterified fatty acid (NEFA) secretion profiles from adipocytes [49]. (**B**) Average NEFA secretion profiles during basal and isoproterenol/forskolin stimulation of adipocytes using the chip in (A) for stimulation and sampling. (**C**) On-chip culture systems for primary adipose tissue [42]. Primary mouse adipocytes were cultured in a 3D collagen matrix (top left) on an 8-channel microfluidic sampling device. Alternatively, explants were trapped using customized traps. 3D CAD rendering (top right) is shown with 3D-printed explant traps (middle right). (**D**) Temporal program of combined insulin, glucose, and fatty acid inputs to adipose explants using a microfluidic multiplexer (μMUX) device [39]. (**E**) Real-time fatty acid uptake responses from treatments in (D) showed insulin-dependent exchange rates. (**F**) Representative fluorescent images of on-chip explants used to generate data in (E). **(G)** Compiled data from (E) showed insulin-dependent fatty acid exchange dynamics. (**H**) Assignment map of the programmable, automated μMUX device, showing the 10 required input channels and one output waste channel used for (D)–(G). Reproduced

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Table 1

List of important adipokines in WAT*

TNF-a tumor necrosis factor alpha, IL-6 interleukin 6, IL-18 interleukin 18, RBP4 retinol binding protein 4, CCL2 CC-chemokine ligand 2, Nampt nicotinamide phosphoribosyltransferase (also called pre-B-cell colony-enhancing factor 1, PBEF1, or visfatin), ANGPTL2 angiopoietin Like Protein 2, CXCL5 signifies one of the cytokine-like proteins with a highly conserved ELR motif preceding the N-terminal cysteine, SFRP5 secreted frizzled-related protein 5

* List modified from Ouchi et al. [19]

Table 2

List of important batokines in BAT

NGF nerve growth factor, VEGFA vascular endothelial growth factor A, BMP8B Bone morphogenic protein 8b, FGF21 fibroblast growth factor 21, L-PGDS Lipocalin-type prostaglandin D synthase, IGF-1 insulin-like growth factor 1, IGFBP-2 insulin-like growth factor binding protein 2, Metrnl meteorin-like, NRG4 neuregulin 4, sLR11 soluble form of the low-density lipoprotein receptor relative, Slit2-C slit homolog 2 protein, C-fragment, PM20D1 peptidase M20 domain containing 1

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Table 3

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adipocyte perfusion.

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