

# Modelos experimentales para estudiar el tejido adiposo en obesidad infantil

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## Experimental models to study adipose tissue in childhood obesity

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# Abstract

Obesity is a major public health concern that has significant repercussions on individuals' well-being and healthcare systems. It is associated with a reduced lifespan and is a risk factor for several chronic diseases, such as type 2 diabetes, coronary heart diseases, and certain

- 5 types of cancers. The understanding of obesity's underlying physiological mechanisms remains limited, despite its significant impact. Childhood obesity is increasingly prevalent and, like adult obesity, it is closely linked to various metabolic diseases. Given the growing occurrence of obesity in children and its consequential clinical and physiological implications, there is a pressing need for targeted experimental research. It is crucial to employ specific cell and animal models to unravel the molecular and genetic factors that
- contribute to obesity in this age group. Here, we review the emerging field of childhood obesity, focusing on the different types of adipose tissue, differences between adipose tissue in adults and children, current available experimental models to study adipose tissue in children, and factors involved in the pathogenesis of childhood obesity.
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Keywords: Obesity, Adipose tissue, Children, Experimental models

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### **Childhood obesity**

Obesity is a global threat. While it was previously predominant in adults, obesity is becoming more prevalent in younger generations, including children younger than 10 years old. In 2019, the World Health Organization (WHO) estimated there were 38.2 million children

under the age of 5 years suffering from overweight or obesity. Data from 2016 revealed that over 340 million children and adolescents (18%) aged 5–19 years were obese or overweight. Furthermore, the WHO also reported that, while overweight and obesity were previously only considered a concern in high-income countries, they are exponentially growing in both low and middle-income countries, with a marked emphasis on urban areas (2019)<sup>1</sup>. Thus, efforts
should be made at the research, clinical, and social levels to stop the alarming growth of

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obesity and its associated metabolic diseases both in adults and children.

# Adipose tissue: colours and functions

Adipose tissue (AT) is a complex and highly dynamic endocrine organ with intricate
metabolic activity that greatly influences the function of nearly all other organ systems through its various adipokines, lipokines, lipids, and peptides<sup>2</sup>. The distribution, localization, and colour of AT determines its function. AT beneath the skin is known as subcutaneous adipose tissue (SAT), whereas AT surrounding internal organs is known as visceral adipose tissue (VAT)<sup>3</sup>. In mammals, there are four colours of AT: white, brown, beige, and pink. In
brief, the major function of white adipose tissue (WAT) is to store unused energy in the form of triglycerides and release it as fatty acids in times of systemic energy need, whereas brown adipose tissue (BAT) uses fuels, such as glucose or lipid, to maintain the core body temperature and function as a thermogenic tissue<sup>4</sup>. Beige adipocytes arise within WAT either due to differentiation of white adipocyte precursors or trans-differentiation of already mature
white adipocytes. Beige and brown adipocytes share many functional and morphological

traits. They take up glucose and triglycerides to generate heat in response to cold exposure and exercise<sup>5</sup>. Finally, pink adipocytes are alveolar epithelial cells within the mammary gland that are responsible for producing and secreting milk<sup>6</sup>. Several models have been used to study adipose tissue in adults and children: 1) *in vitro* cell cultured lines and primary adipocytes derived from both animals and humans and 2) *in vivo* animal models (summarized

5 adipocy

# in Table 1 and section "Models to study AT" below).

### Differences in AT between children and adults

- The characteristics of AT and its response to pathological inputs, like obesity, differ between children and adults<sup>7,8</sup>. In general, AT growth can be classified as either hypertrophic, in which 10 the adipocyte size increases, or hyperplastic, in which the number of adipocytes increases<sup>9</sup>. Body fat mass expands significantly due to cell size enlargement (hypertrophy) up to 6 months of age. Nonetheless, from the age 6 months until early adolescence (around 10 years old), the fat content increases alongside cell number<sup>10,11</sup>. Puberty (10–18 years old) is 15 characterized by an increase in both hypertrophy and hyperplasia. This stage is also marked by sexual dimorphism. Indeed, sexual dimorphism typically emerges during embryonic stages. However, in the case of obesity, it is during puberty that these differences become significantly pronounced and continue into adulthood. Therefore, the utilization of cell and animal experimental models becomes crucial in investigating this important aspect. 20 Subcutaneous fat distribution differs between males and females. Around the age of 18 years, under normal weight conditions, cell number reaches a plateau, establishing the adipocyte number for the future<sup>10,11</sup>. In adulthood, the capacity for preadipocyte differentiation declines, and therefore, fat content increases mainly due to adipocyte hypertrophy, in contrast to childhood and adolescence<sup>12,13</sup>. In addition, aging has been shown to foster AT redistribution
- 25 by significantly reducing SAT and simultaneously increasing VAT<sup>13</sup>. This rearrangement has

been strongly suggested to explain why elderly people display a worse metabolic health than younger generations<sup>13</sup>.

### Models to study AT

- 5 Studies on childhood obesity, along with its global prevalence, is a relatively recent phenomenon and, consequently, data from cell and animal models are sparse. A deeper understanding of the molecular mechanisms regulating adipocyte function and disfunction in children is necessary to plan strategies to treat and prevent obesity and related metabolic complications. Over the past few decades, several preadipocyte and mature adipocyte cell
- 10 models have been used to study adipogenesis in adults. Rodent cells lines are the most widely used cell line models, although both porcine and feline primary cells have also been used to a lesser extent<sup>14,15</sup>. Among rodents, mouse preadipocytes including 3T3-L1, 3T3-F442A, OP9, and C3H10T1/2 stromal cells have been studied in detail over the years, with 3T3-L1 being the most established and commonly used cell line<sup>15</sup>. However, porcine primary preadipocytes
- 15 have been suggested as a better model for obesity research due to their greater similarity to human cells<sup>15</sup> (Table 1).

In addition, human cell lines (ChubS7<sup>16</sup> and imAPAD<sup>17</sup> and imGPAD<sup>17</sup>) and primary cells have been increasingly used to study adipocyte biology and obesity-related changes. However, only one preadipocyte cell line, Simpson-Golabi-Behmel-Syndrome (SGBS), has

- 20 been isolated from children<sup>18</sup>. This cell line was derived from the AT of an infant with SGBS, which is a quite rare syndrome characterized by macroglossia, macrosomia, renal and skeletal abnormalities, and an increased risk of embryonal cancer, distinct from polygenic or monogenic obesity. While the insights gained from studying SGBS cells are undoubtedly valuable, it is important to consider that the results may be applicable only within a specific
- 25 context. A recent study has shown that these cells are metabolically distinct from primary

adipocytes<sup>19</sup>, indicating the importance of establishing child-derived cell models from healthy

and obese children for further studies (Table 1).

**Table 1**: Models to study adipose tissue.Summary of adipose in vitroand in vivo models discussed in this paper

Animal cell line models <sup>14</sup>	
Mouse 3T3-L1	Derived from 17- to 19-day-old Swiss 3T3 mouse embryonic fibroblasts, most used cell line model to study adipogenesis, high number of passages
Mouse 3T3-F442A	Derived from murine Swiss 3T3 cells. Able to accumulate higher fat content than 3T3-L1
Mouse OP9	Derived from the calvaria of new-born mice. Fast adipogenic differentiation (72 h). Suitable for high-throughput assays
Mouse C3H10T1/2	Derived from 14-17-day-old C3H embryonic stem cell precursors. Frequently used to study adipogenesis
Primary animal cells <sup>13,14</sup>	
Porcine cells	Better model for adipogenesis and adipose tissue-related complications compared to rodents due to their higher similarity towards human cells
Feline cells	Used to study adipogenesis and type 2 diabetes
Human cell line	models
ChubS7 <sup>15</sup>	Derived from abdominal subcutaneous adipose tissue of a 33-year old obese female, high proliferation and differentiation rate
imAPAD <sup>16</sup>	Derived from abdominal subcutaneous adipose tissue of a 50-year old healthy male, high proliferation and differentiation rate. Retain depot-specific function
imGPAD <sup>16</sup>	Derived from gluteal subcutaneous adipose tissue of a 50-year old healthy male's, high proliferation and differentiation rate. Retain depot-specific function

SGBS <sup>17</sup>	Derived from subcutaneous adipose tissue of a 3-month-old infant with Simpson-Golabi- Behmel syndrome. Used to study adipogenesis. Metabolically distinct from primary adipocytes
Primary human cells	
Primary Preadipocytes <sup>14</sup>	Primary cells, represent the most physiological model of adipose precursors. Retain their depot-of-origin and individual traits. Limited number of passages, not always easily accessible
<i>In vivo</i> animal Models	
C. elegans <sup>19</sup>	Used to study factors involved in food satiety
D. melanogaster <sup>19</sup>	Used to study neuronal pathways implicated in fat deposition
<b>D. rerio</b> <sup>19</sup>	Frequently used model for the study of adipose tissue biology and obesity-related diseases
Zucker rat pups <sup>20</sup>	Used to study AT growth from suckling to puberty stages
Small litter size rodents <sup>23</sup>	Better model to study childhood adiposity andlateronset ofmetabolic diseases than large litter size

*In vivo* models are also crucial for assessing both basic and clinical assays. Small animal models, such as *Caenorhabditis elegans*, *Drosophila melanogaster*, and *Danio rerio*, have displayed many advantages due to their short life cycle, known genome maps, and low maintenance cost, for studying the mechanistic underpinnings of various pathological conditions. Obesity research in *C. elegans* has uncovered factors involved in both food sensing and food satiety. Conversely, studies using *D. melanogaster* identified neuronal pathways involved in fat deposition<sup>20</sup>. While small animal models can be valuable for various preclinical studies, they may not be suitable for evaluating the multiorgan system effects of obesity in adults or children. Nevertheless, these animal models can contribute to

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understanding the molecular and neurological pathways that may be influenced or disrupted

by childhood obesity. Hence, small rodents, such as mice and rats, are widely employed as mammalian *in vivo* model systems for obesity and related disorders at the genetic, molecular, and functional levels<sup>20</sup>. Zucker rat pups have been used to study AT growth from suckling to puberty stages. Use of this model revealed that the highest rates of adipocyte proliferation, *de* 

5 *novo* lipogenesis, and outbreak of excessive fat mass occurred at the end of the suckling stage (Table 1)<sup>21</sup>. The premise that human metabolism and physiology differ to that of animals and that adults AT development is different between children and adults limits the insight that these studies may provide on obesity and related illnesses in children<sup>14,15</sup>.

### 10 Factors involved in the pathogenesis of childhood obesity

In terms of socioeconomic and environmental factors, a child's body mass index (BMI) and its mother's BMI and education status have been reported to be associated with risk of becoming obese<sup>22</sup>. These three risk factors are particularly noteworthy from the age of 6 years onwards. Overweight and obesity were markedly prevalent among children who previously

15 displayed a cluster of all three, providing both easy and quick evaluations and further prevention assessments at clinical encounters<sup>22</sup>.

Furthermore, some studies have attempted to uncover epigenetic patterns directly linked to emerging childhood obesity. These transcriptional modifications may offer insight into predisposition as they switch in response to environmental factors, such as physical activity

- 20 or nutrition, and differ between different ethnicities. An example of this is the epigenetic modification of hypothalamic feeding circuits in offspring when subjected to stress during pregnancy or postnatal periods. This alteration leads to changes in the expression of neuropeptides, affecting their regulation of feeding behaviour<sup>23, 24</sup>. As a result, energy metabolism is altered in the offspring increasing their susceptibility to develop obesity and
- 25 metabolic-like syndromes in adulthood<sup>23</sup>. These epigenetic modifications include mutations

in gene promoters associated with both appetite regulation and stress. Additionally, adverse experiences during early childhood can lead to DNA methylations in genes associated with obesity<sup>23,25</sup>.

DNA methylation of the nuclear respiratory factor 1 (NRF1) gene at a crucial CpG
dinucleotide was strongly correlated with obesity emergence at age 3-5-years in nonobese Hispanic children. However, data on other ethnicities has not been reported<sup>26</sup>.

While the influence of mothers on the transmission of obesity risk to their offspring has been extensively studied<sup>23,24,27,28</sup>, only few studies have evaluated the potential contribution of fathers to their children's obesity development<sup>27,29</sup>. However, recent investigations indicate

- 10 that interventions aimed at reducing obesity in men, such as exercise and weight loss, can have a positive impact on sperm quality and alter patterns of DNA methylation in sperm<sup>27</sup>. In order to gain a comprehensive understanding of the underlying intergenerational mechanisms involved in early human exposure, it is crucial for future research to address both maternal and paternal epigenetic and epidemiological studies. Studying all these factors in detail will benefit from the establishment of reliable models to study childhood obesity, including
- child-derived cell models, and animal models.

### **Conclusions and future perspectives**

Obesity and its associated metabolic diseases, such as diabetes, insulin resistance, cardiovascular disease, asthma, and some types of cancer, are becoming more prevalent worldwide. This is especially worrisome at younger ages. Importantly, research into obesity has largely focused on adults and overlooked in children. *In vivo* and *in vitro* models to study childhood obesity are still very limited. Thus, experimental models to study AT in childhood obesity are required to elucidate accurate early prediction markers and, ultimately, appropriate prevention strategies.

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