

Obesity Leads to Elevated Level of Circulating Cell-Free DNA



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Abstract

With the steadily growing number of studies in the issues related to obesity research, it is proven, that obesity might be related to wide number of body disorders, some of these disorders have been already studied and investigated, and some still needed more investigations, which has opened new horizons to unlimite hypotheses about involvement of obesity in many physiological disorders. In the current opinion, we hypothesized to the conclusion that there is a correlation between obesity severity and elevated cfDNA levels in the circulation. This correlation is based on free radicals and oxidative stress hypothesis of cellular and molecular damage in cell injury, which in turn leads to a disorder in the whole body and enhances the inflammatory status.

Keywords: Obesity; Inflammation; Oxidative stress; Free radicals; Cfdna

Abbreviations: WHO: World Health Organization; NIH: National Institutes of Health; BMI: Body Mass Index; LEPR: Leptin Receptor; LEP: Leptin; POMC: Pro-Opiomelanocortin; MC4R: Melanocortin-4 Receptor; SNP: Single-Nucleotide Polymorphisms; ROS: Reactive Oxygen Species

Introduction

It is clear to the medical community the importance and impact of the various types of obesity on a wide range of diseases and physical disorders and requires more research and investigation in any relationship suspected between obesity and other disorders. This review emphasizes a new opinion and summarises the current state of knowledge about the impact of obesity on level of cell-free DNA in circulation, which has been used recently as a new and significant biomarker for some illness and inflammatory disorders.

Obesity has received significant attention as a complex health hazard. Obesity is defined as a status of excessive fat accumulation in adipose tissue which is resulting in health is impairment [1]. The National Institutes of health (NIH) and The World Health Organization (WHO) have used Body Mass Index (BMI) which is the most widely used measure of obesity due to its low cost and simplicity. BMI is defined as the weight in kilograms divided by the height in meters squared, according to BMI overweight means as a BMI between 25.0 and 29.9 kg/m²; and obesity as a BMI greater than 30.0 kg/m² [2,3].

The most common methods used as reference measures of body composition especially for research purposes include single-cut imaging of the abdomen using computed tomography scan or magnetic resonance imaging, densitometry, and dual-

energy X-ray absorptiometry. However, increasing evidence indicates that abdominal obesity is also useful beside total body fat, and it is independent predictor of several cancer and cardiovascular-related outcomes in addition to other illnesses [3]. Some of the commonly used measures of abdominal obesity are hip circumference, waist circumference, and waist-to-hip ratio [2,3].

Prevalence of Obesity

Obesity is a complex, multifactorial, and one of the greatest public health epidemics of the 21st century [4]. The prevalence of obesity is rising globally, with about 2 billion adults worldwide currently classified as being overweight or obese [5]. However, estimates of its prevalence are still insufficient for all countries, and the available data are not comparable or uniformly accurate [6]. If secular trends continue, by 2030 an estimated 20% will be obese and another 38% of the world's adult population will be overweight [7]. In the USA, based on earlier secular trends point, by 2030 an estimated 85% of adults will be obese or overweight [8]. Whereas in most developed countries growth, trends in overall obesity seem to have levelled off [9], but still morbid obesity in many of these countries continues to rise, including among children. Furthermore, obesity prevalence in most developing countries continues to trend upwards toward US levels [8].

Common Causes of Obesity

Environmental factors are seeming to be major contributors to the obesity. It is certain that obesity evolves when there is an imbalance between increase energy intake and decrease energy expenditure. Evidence supports the contribution of both excess energy intake and decreased energy expenditure in the obesity epidemic [10-12]. On the other hand, it is known that single gene mutations are associated with rare forms of monogenic obesity (leptin receptor (LEPR), leptin (LEP), pro-opiomelanocortin (POMC), and melanocortin-4 receptor (MC4R)) [13]. However, there is growing evidence that single-nucleotide polymorphisms (SNP) or common genetic variants may play an important role in the obesity. These SNPs have some effects on an individual susceptibility to obesity, but due to their high frequencies, if involved in lipogenic pathways, may be able to have a wide contribution to obesity in the population [14]. Besides, nutrigenetics plays a vital role [15]. The methods used for analysis these mutations are limited and there is a need to work with a cos-effective method [16,17] and to consider any particular method as a prominent bimolecular tool for obesity and its severity.

Obesity and Oxidative Stress

It has been reported that obesity is associated with chronic systemic inflammation in adipose tissue. This condition is affected by the activation of the innate immune system in adipose tissue that enhances oxidative stress and pro-inflammatory status, affecting a systemic acute-phase response. Various chronic diseases are also the result of obesity (e.g., diabetes mellitus, metabolic syndrome, cancers, and cardiovascular and liver diseases) and associated with oxidative stress [18]. So, it has been hypothesized that inflammation of adipose tissue in obese patients plays a significant role in the pathogenesis of obesity-related disorders [19].

Adipose tissue is a storage organ required for energy homeostasis. Adipose tissue is composed of adipocytes primarily and contains other cells (e.g. pre-adipocytes, endothelial, fibroblasts, fibroblastic and immune cells) [20], secreting hormones and cytokines (adipocytokines or adipokines) which exercise autocrine, endocrine, and paracrine action on the whole body. In pathological and, even more, in physiological conditions, adipokines also induce the production of reactive oxygen species (ROS), generating oxidative stress and, in turn, a major, irregular output of other adipokines [21]. Oxidative stress and pro-inflammatory processes are strongly associated [22,23]. Upon activation, many immune cells produce free radicals, and, in the same way, the synthesis of ROS enhances the inflammatory status.

Oxidative Stress and Cell Aging

Oxidative stress is also considered as a measure of unbalance between the production of free radicals and antioxidant defences [24-26]. Oxidative stress has been involved in response to stress

and the pathogenesis of psychiatric and neurologic diseases [27-29]. Mostly, the main cause of oxidative stress has been a production of free radicals. Free radicals are chemical species which contain an odd number of electrons in last electronic layer [30]. This state of electrons of the last layer that grant high reactivity to these molecules or atoms [31]. ROS often generated by mitochondria, in addition to other sources of ROS, such as NADPH oxidase (NOX) enzymes; they are a group of membrane proteins with a known function to generate ROS. These enzymes are working as a transmembrane electron transport chain by using cytoplasmic NADPH as an electron granter to O₂ molecule to generate superoxide anion in the lumen of intracellular organelles or the extracellular space. Superoxide anion is mostly considered the essential product of the electron transfer, also to generate other ROS in particular hydrogen peroxide [32,33].

The researchers have identified a total of seven NOX genes: NOX1 to 5 and DUOX1 and 2. The better-described isoform NOX2 needs interaction with another trans-membrane protein, p22phox, in addition to the cytosolic subunits, p40phox, p47phox, p67phox, and with one of the small Rho GTP-binding proteins, Rac1 or 2. On the other hand, other NOX isoforms have a different mechanism of activation and require p22phox for activity. NOX3 and NOX1 enzymes require interaction with cytosolic subunits, NOXO1 and NOXA1, and with Rac1 or 2 [34]. NOX4 seems to be constitutively active; NOX5 and DUOX enzymes are often regulated by increased intracellular Ca²⁺ [33]. It is worth mentioning that ROS generated by NOX enzymes can directly impact cellular functions; by encourage the oxidation of proteins, then their functional and structural changes [13]. From this particular issue we suggested that increases oxidative stress which is enhances the inflammatory status, subsequently causes damage and destruction of cells which is increasing cell-free DNA in circulation.

Circulating Cell-Free DNA

In 1948, Mandel and Metais discovered the presence of cell-free DNA (cfDNA) in blood plasma [35]. This discovery was not attracted great attention in the medical scientific community until 1994; when the importance of cfDNA was explored in the study of mutated RAS gene fragments in the circulation of cancer patients [36,37]. Understanding the mechanisms of generation of cfDNA is important for conclusion its role in pathology and biology and speeding the translation of analyses to clinical practice. Despite the apparent of cfDNA in blood circulation and other bio-fluids, the exposition of its origin has only been a partial triumph [36,37].

Mostly, cfDNA are found as double-stranded molecules with molecular weights in the wide range of 0.18 kB to 21 kB [38,39]. Various possible sources and cognate mechanisms have been detected. Firstly, it was supposed that cfDNA enters the blood following the cells lysis on the interface between a tumour and blood circulation only. Since it was reported in patients that the concentration of cfDNA in their blood is higher than could be

accounted for by the mass of cells present, this hypothesis has been abandoned [40]. The exact mechanism of releasing cfDNA into the blood circulation is still unknown, several researchers have suggested that in either healthy or diseased individual's cfDNA is predominantly haematopoietic origin [41,42]. They also demonstrated that under normal conditions, cfDNA is released from apoptotic rather than necrotic processes. It should be emphasized on the role of a dynamic balance between processes of cellular DNA secretion/release and mechanisms of DNA degradation and clearance in the keep of the existing level of cfDNA in human blood [43,44]. The cfDNA is composed of both genomic DNA (gDNA) and mitochondrial DNA (mtDNA) [45]. The concentration of cfDNA has been assessed previously by quantitative PCR (such as TaqMan and SYBR Green) or by different fluorescence-based methods (such as, ultraviolet (UV) spectrometry and PicoGreen staining) [46].

Based on the above discussion, increased oxidative stress, which might result in enhancing inflammatory status and result in cell damage can therefore proposed to rise in circulation of cfDNA. Therefore, various metabolic diseases also can have high chances of high cfDNA level in circulation according to hypothesis of related cfDNA with oxidative stress. The significant disorders might be associated with cfDNA are metabolic diseases high oxidative stress, for example, obesity and diabetes [47-49]. Studies are required in future to look into the effect of severity of metabolomics stress and oxidative stress on cfDNA [50].

Conclusion

Looking at the cited literature, we hypothesized to the conclusion that there is a correlation between the obesity and elevated cfDNA levels. This correlation based on free radicals and oxidative stress hypothesis of cellular molecules damage and cell injury which in turn leads to a disorder in the whole body, but it still needs to more study and investigation.

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