



Review

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Immunometabolic Links Underlying the Infectobesity with Persistent Viral Infections

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ABSTRACT

Obesity and its related comorbidities are prevailing globally. Multiple factors are etiological to cause obesity and relevant metabolic disorders. In this regard, some pathogenic infections including those by viruses have also been associated with obesity (termed especially as infectobesity). In this mini-review, I examined recent publications about primary or cofactorial role of viral infections to exacerbate the local and systemic immunometabolic cues that underlie most cofactorial obesity. Major immuno-metabolic pathways involved, including that mediated by interferon (IFN) signaling and peroxisome proliferator activated receptor- γ (PPAR- γ), are discussed.

Introduction

Obesity manifests as metabolic overload of excess fat in adipose depots, but entails various immunological disorders. This can be further worsen the overweight into a metabolic syndrome as well as other life-threatening complications including diabetes, heart disease, liver steatosis, and cancer^{1,2}. Obesity and its related comorbidities are epidemic globally to reach an alarming level. Various factors from predisposing genetics to habitat environment could contribute to the complex disorders relevant to obesity epidemic¹⁻³. Despite intensive studies, the etiological keys that cause metabolic disorders and obesity remain incompletely understood³. From an evolutionary view, animal metabolic and immune systems integrate as the most fundamental requirement for a species survival⁴. In this mini-review, I will focus on the obesity induced by energy-dense diets and viral agents to examine the major immunological links that drive relevant metabolic disorders and obesity development⁵.

Viral Infectobesity

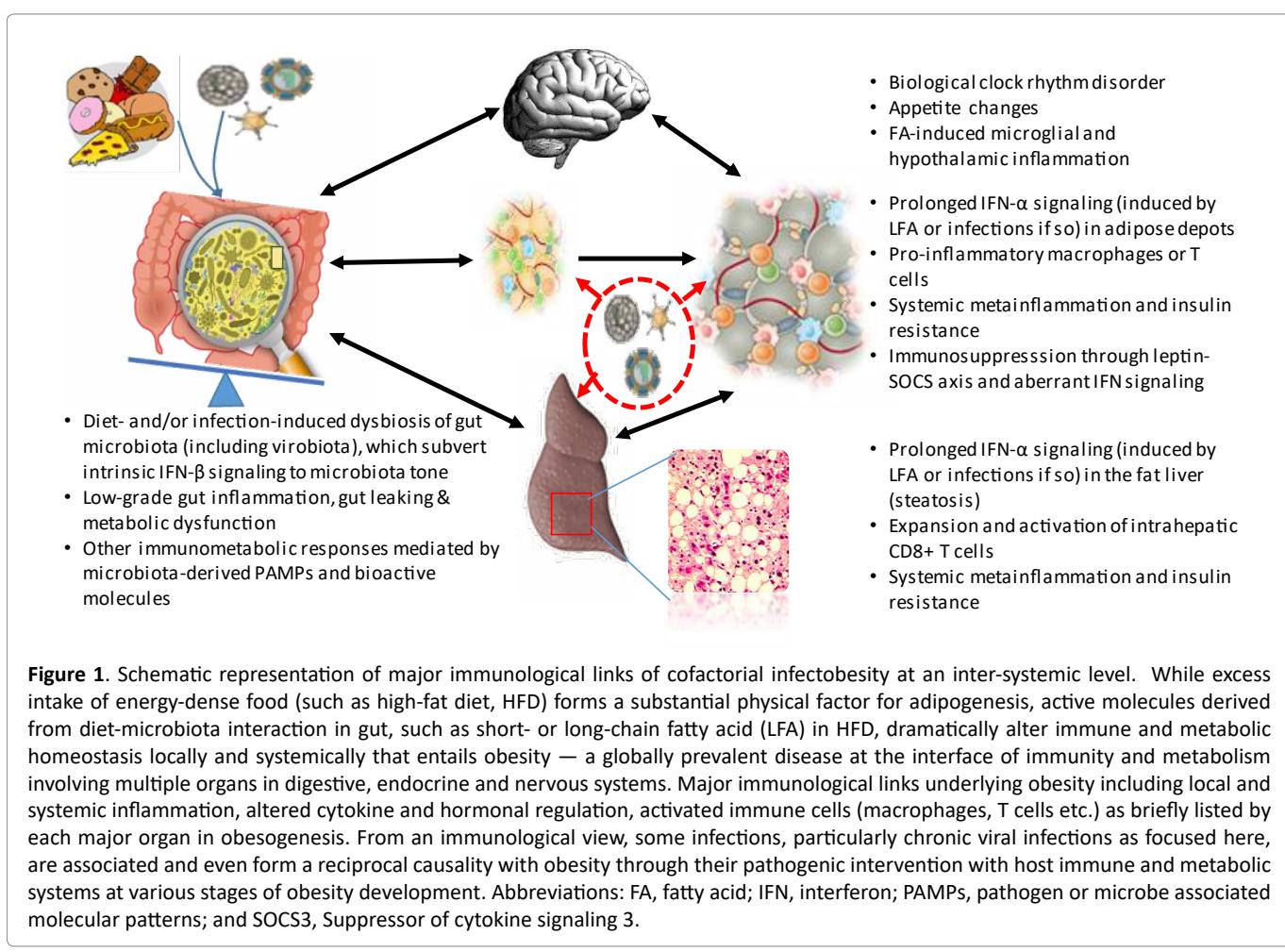
Infectobesity describes the association of microbial infections with obesity and its relevant morbidities, which has been often observed in humans and reproduced in animal models⁵⁻⁸. The major viruses and virus-like agents (VLA) that have been associated with obesity include members in adenoviridae, herpesviridae, phages, transmissible spongiform encephalopathies (slow virus), and hepatitides⁵⁻⁸. For example, at least five human adenovirus (Ad) serotypes, i.e. Ad-5, -9, -31, -36, and -37, have been associated with obesity at different levels. These adenoviruses demonstrated activity to induce adipocyte differentiation and increase lipid synthesis in cultured animal fibroblasts or pre-adipocytes. Serotypes Ad-5, Ad-37, and particularly Ad-36 infections led to

the increase of adipogenic activity in animal models of chickens (Ad-36 and -37), mice (Ad-5 and -36), rats and marmosets (Ad-36)⁵⁻⁸. Cytomegalovirus (CMV) as a common herpesvirus that infects over half of adults by age 40, was associated with metabolic syndrome (diagnosed as systemic inflammation, insulin resistance and gut barrier disruption etc.) particularly in females⁹. Recent additions to this list also include Dengue fever virus (DEGV) and treated-HIV infections (which belong to Flaviviridae and retroviridae, respectively) in the patients post the viral infections entering into a latent phase, such as after antiretroviral therapy (ART) in HIV cases¹⁰⁻¹³. Notably, HIV infection itself may not cause lipohypertrophy; however, metabolic syndrome and visceral adiposity have been observed in treated patients probably due to the viral persistence and chronic inflammation resulted from antiretroviral therapy^{12,13}. Owing to recent advancement in viral metagenomics studies (virome), it is expected that more previously unknown viruses may be identified to be “pawns”, which defines a responsible taxon of microbiota that subjects to adipogenic regulation. The increasing pawns may partly result from the selection pressure of current obesity-prone life styles in people and contemporary fattening management in livestock^{5,14-17}. It

is remarkable that most obesity-associated viruses have a livestock reservoir or share similar pathogenic persistence between livestock and people^{5,15-17}.

Major immunological links underlying the viral infectobesity

Figure 1 illustrates most immunological links underlying obesogenesis but not limited to viral infectobesity. A majority of obesity cases are developed owing to gradual cofactorial induction that builds up immunometabolic disorders involving multiple organs in digestive (including gut microbiota), adipose, endocrine and nervous systems¹⁸⁻²¹. In the situation of viral infectobesity, excess intake of energy-dense food (such as high-fat diet, HFD) still forms a substantial physical factor for adipogenesis. Critically, active molecules, such as short- or long-chain fatty acid (LFA) that derive from diet-microbiota interaction in gut, dramatically alter immune and metabolic homeostasis locally and systemically²²⁻²⁴. This further leads to obesity and relevant comorbidities in involved organs (Figure 1)^{21,24}. Major immunological links underlying obesity include: (1) local and systemic inflammation across intestine, adipose tissues, liver and even hypothalamus in the brain, (2) altered cytokine (such as TNF- α and IFNs emphasized



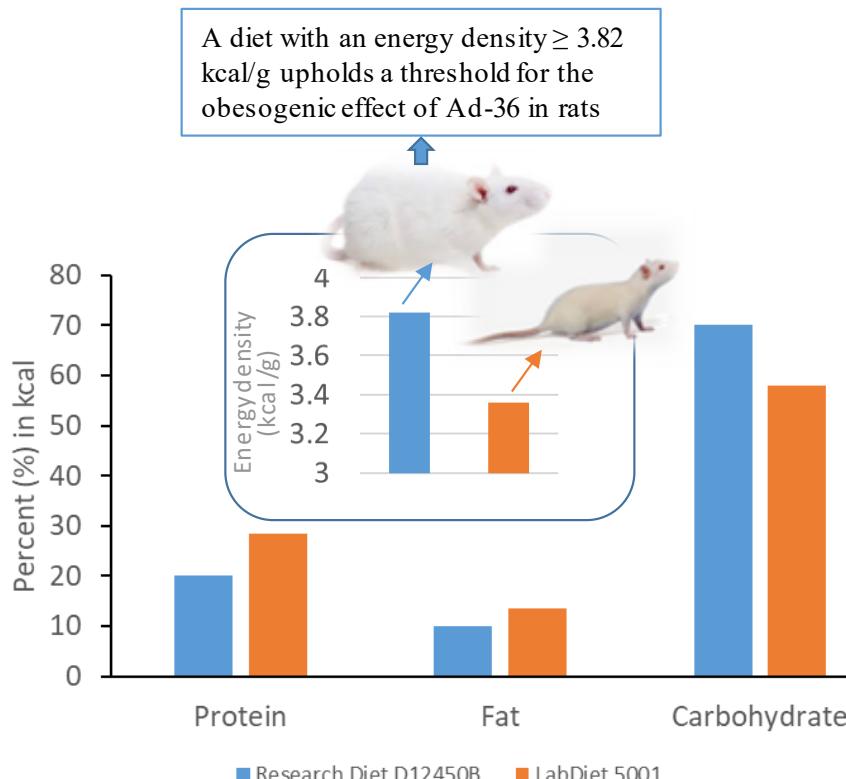


Figure 2. The high energy density of a diet upholds the viral effect in induction of viral infectobesity. Comparison of caloric information and energy density (inlet framed) of two diets that showed different synergic effect with Ad-36 in obesity induction as in two studies, respectively^{45,46}. As both diets have similar amount of 10% fat in kcal, the Research Diet D12450B, which showed synergistic effect with Ad-36 for obesity induction, contains about 12% higher carbohydrate in kcal but 8.5% lower protein in kcal than the LabDiet5001 that showed no synergistic effect. Overall, the energy density of the D12450B diet is about 0.5 kcal/g higher (3.82 vs. 3.36) than the LabDiet 5001. The diet formulas and contents were directly extracted and calculated from the producers' websites at <https://researchdiets.com/formulas/d12450b> and <https://www.labdiet.com/Products/StandardDiets/index.html>.

here) and hormonal regulation, and (3) improperly activated immune cells (such as adipose macrophages and intrahepatic T cells)²¹⁻²⁵. From an immunological view, some infections, particularly persistent viral infections as focused here, are associated and even form a reciprocal causality with obesity through their pathogenic intervention with host immune and metabolic systems at various stages of obesity development¹⁹⁻²⁵. In this context, we stress the regulatory role of interferon (IFN) system in this virus-involved adipogenic process because recent studies have highlighted it as an immunological axis in the obesogenesis of viral infectobesity²⁵⁻³⁰. First, an intrinsic IFN- β signaling response to gut microbiota tone was demonstrated to be critical to maintain gut homeostasis³¹. A microbiota dysbiosis caused by HFD and persistent viral infections may subvert this intrinsic IFN signaling and progressively trigger low-grade gut inflammation, gut leaking and increase caloric intake and storage in both intestinal and adipose depots^{24,25}. Second, persistent viral infections and LFA produced during adipogenesis irritate a prolonged IFN- α signaling, which has been detected in both adipose depots and fat liver during obesity development

to be a biomarker for persistent viral infections and some autoimmune diseases²⁴⁻³⁰. Third, the prolonged IFN signaling contributes to the immune activation of pro-inflammatory macrophages and cytotoxic CD8+ T cells in adipose tissues and fat liver, which represent major cell markers for obesity and metabolic syndrome²⁵. In contrast to induce effective antiviral immunity, the prolonged IFN signaling actually causes an immune dysfunction against further viral infections and aggregates to a relevant immune suppression induced by adipose cytokines (such as leptin through SOCS3). This immune deviation may form a reciprocal causality between virus susceptibility and obesity resulting in the complication of viral infectobesity^{7,27,30-35}. In addition, chronic inflammation and immune senescence accompanying aging process generally coincide with a high incidence of viral infections in senior people^{36,37}. It has been observed that a 5-8% higher rate of obesity in peoples of the age groups over 40 and 60 years old^{1,2}. The potential correlation of the higher occurrence of viral infections and obesity rates in the aged groups ascribes another big topic in viral infectobesity and warrants further investigations^{1-3,36,37}.

The metabolic basis upholding viral infectobesity

It has a chicken-or-egg disputation about infectobesity. Among a dozen of aforementioned viruses, most of them are associated with obesity and relevant morbidities through epidemiological statistics and viral coexistence (or seropositive to viral antigens) during the obesogenesis in humans and animals⁵⁻¹³. However, few of them have been applied to the complete tests per Koch's postulates to establish as obesogenic pathogens³⁸. In this context, human adenoviruses (Ad), particularly serotype Ad-36 have been well characterized in their obesogenic effect in both cell and animal models including chickens and rats^{5,39}. The adenoviral E4orf1 protein (early region 4 open reading frame 1) was identified to be a viral mechanism mediating acute adipogenesis in both cells and animals^{40,41}. Indeed, transfection of E4orf1 gene alone mimicked Ad-36's effect on the increase of glucose uptake in preadipocytes, adipocytes, or myoblasts, but reduced glucose output by hepatocytes⁴⁰. Functionally acting through the Ras/PI3K signaling pathway, E4orf1 protein actually improved hyperglycemia and insulin disposal in vitro, implying a potential for anti-diabetic therapy development^{40,41}. Peroxisome proliferator activated receptor-γ (PPAR γ) is so-called a master-switch of adipocyte development. Ad-36 was capable of inducing the expression of cellular glucose transporters to facilitate energy supply and the activation of PPAR γ to signal adult stem cells differentiating into adipocytes⁴²⁻⁴⁴. Despite the positive correlation described above, the obesogenic effect of Ad-36 was not well reproduced in mouse and monkey models^{5,39}. Similarly, the association of Ad-36 infection with human obesity was not always significant when tested across different subject cohorts^{5,39}. In our recent study using a rat model⁴⁵, we showed that not Ad-36 alone but co-administration of Ad-36 and high-fat diet (HFD + Ad-36) induced significant obesity in rats at 8 weeks after induction. To determine what caused the difference between a previous report and ours in the obesogenic effect of Ad-36 in rat models^{45,46}, we compared the caloric information and energy density of two diets, which were used in a previous report and our study to show difference in synergistic effect with Ad-36 in obesity induction, respectively^{45,46}. As both diets have similar amounts of fat at ~10% in kcal, the Research Diet D12450B (Research Diets, New Brunswick, NJ) used in previous study⁴⁶ that facilitated Ad-36 for obesity induction contains about 12% higher carbohydrate in kcal but 8.5% lower protein in kcal than the LabDiet5001 used in our study⁴⁵. Overall, the energy density of the D12450B diet is about 0.5 kcal/g higher (3.82 vs. 3.36) than the LabDiet 5001 (Figure 2)^{45,46}. In addition, the other components including minerals, vitamins, and fiber are similar between the two diets. Thus, this comparison implies that both the fat content and the energy density (contributed also by the other diet components, especially

carbohydrate) of a diet, should be considered to uphold Ad-36 effect in obesogenic induction. This is important in terms of determining co-factorial interaction in obesity induction, indicating a certain level of energy density in a diet stands as a basis for cofactorial infectobesity at least for Ad-36 (Figure 2)⁴⁵⁻⁴⁷. Collectively, adiposity induced by energy-dense diets may set up a metabolic aberration that facilitates the virus obesogenic effect; or vice versa viruses may snatch the cellular metabolic status to benefit viral replication and exacerbates adipogenesis at the early stage of the viral infectobesity^{5-8,45-47}. However, at the mid and late stages of obesity development, especially after the manifestation of metabolic syndrome or organ pathologies, the immunological aberration arising from adiposity and particularly persistent viral infections will play more and more role in aggravating obesity and related comorbidities as described in previous Section^{4,46,47}.

Concluding remarks

Due to co-factorial integration in viral infections and obesity, it may be impractical to determine which factor comes first in triggering and manifesting the infectobesity⁵⁻⁸. I reinforce the immunometabolic links that underlying the reciprocal causality of viral infections and obesity in the infectobesity^{4,19-21}. Along with other infectious agents beyond persistent viruses that associated with obesity, a general immunopathological common about pathogenic persistence and relevant immune deviation is a theme underlying the infectobesity^{4,19-21}. In most cases, obesity and viral infections may interact dynamically as a reciprocal causality rather than a simple sequential relationship. Moreover, IFN (particular type I IFNs) responses have recently been postulated as a central immunological axis that governs adipocyte differentiation and T cell pathogenicity during obesity-associated metabolic diseases^{25,48-50}. In summary, immunological dysfunction accompanying persistent viral infections and obesogenesis provides a key for understanding and potential means to treat viral infectobesity^{4,25,48-50}.

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Conflicts of Interest

The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript, or in the decision to publish the results.

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