



The impact of obesity-induced inflammation on clonal hematopoiesis

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Purpose of review

This review meticulously delves into existing literature and recent findings to elucidate the intricate link between obesity and clonal hematopoiesis of indeterminate potential (CHIP) associated clonal hematopoiesis. It aims to enhance our comprehension of this multifaceted association, offering insights into potential avenues for future research and therapeutic interventions.

Recent findings

Recent insights reveal that mutations in CHIP-associated genes are not limited to symptomatic patients but are also present in asymptomatic individuals. This section focuses on the impact of obesity-induced inflammation and fatty bone marrow (FBM) on the development of CHIP-associated diseases. Common comorbidities such as obesity, diabetes, and infection, fostering pro-inflammatory environments, play a pivotal role in the acceleration of these pathologies. Our research underscores a notable association between CHIP and an increased waist-to-hip ratio (WHR), emphasizing the link between obesity and myeloid leukemia. Recent studies highlight a strong correlation between obesity and myeloid leukemias in both children and adults, with increased risks and poorer survival outcomes in overweight individuals.

Summary

We discuss recent insights into how CHIP-associated pathologies respond to obesity-induced inflammation, offering implications for future studies in the intricate field of clonal hematopoiesis.

Keywords

clonal hematopoiesis, clonal hematopoiesis of indeterminate potential, inflammation, obesity

INTRODUCTION

Clonal hematopoiesis of indeterminate potential (CHIP) is linked to increased mortality risk and the development of hematologic malignancies [1,2]. CHIP affects 10–20% of adults over 70 years of age and is due to acquisition of somatic mutations accumulating in hematopoietic stem (HSCs) and progenitor cells (HSC/Ps) [3]. Commonly mutated genes in CHIP include *TET2*, *DNMT3A*, *ASXL1*, and *JAK2*. These mutations provide a competitive advantage to mutant HSCs and HSC/Ps in a process known as clonal hematopoiesis [2,4]. Unlike normal HSCs, CHIP-associated mutations in HSCs lead to skewed differentiation towards myeloid lineage [5]. While mutations in CHIP-associated genes are commonly found in patients with myeloproliferative neoplasms (MPNs) and acute myeloid leukemia (AML), these mutations can also be detected in individuals without any symptoms [6]. It is noteworthy that not all individuals with CHIP-associated mutations develop hematologic malignancies, indicating the presence of predisposing factors that might promote clonal expansion and

disease development. To this end, inflammatory conditions, such as obesity, diabetes, infection, smoking, and pulmonary inflammation, may contribute to CHIP-associated diseases [7–12], suggesting that environmental and additional cell nonautonomous factors likely contribute to clonal hematopoiesis. The exact mechanism(s) are however not yet fully understood. Gaining a better understanding of these mechanism(s) will be crucial in identifying individuals at risk for developing CHIP-associated diseases and stratifying patients for appropriate treatment. Obesity and overweight

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KEY POINTS

- Obesity is associated with preleukemic CHIP mutations and exacerbates clonal hematopoiesis and risk factor for leukemia development.
- Obesity increases accumulation of bone marrow adipocytes, forming FBM. FBM negatively impacts bone mass by impeding osteoblast differentiation, thereby altering hematopoiesis.
- Accumulation of bone marrow adipocytes secrete abnormal amount of pro-inflammatory cytokines such as *IL-6*, *IL-1 β* , and *TNF- α* , contributing to leukemia development.
- CHIP-associated mutations in obese individuals may reprogram adipocytes into cancer associated adipocytes, which subsequently secrete growth, inflammatory, fibrotic and angiogenic factors and lymphokines, contributing to leukemia development.

have been suggested as a risk factor for cancer development [13,14]. However, it is unclear if obesity is also a risk factor for clonal hematopoiesis and for transforming preleukemic stem cells (pre-LSCs) into full-blown leukemia such as AML or severe form of MPN. To this end, we analyzed exome sequencing and clinical data from 47 466 individuals with validated CHIP showed that CHIP is present in number of the study population and is associated with a significant increase in WHR. Importantly, we observed that obesity is associated with myeloid leukemia [15^{***}]. Furthermore, recent studies, alongside our own findings, reveal a strong association between obesity and myeloid leukemias in both children and adults with increased risk and worse survival outcomes in obese and/or overweight individuals [16–20]. Furthermore, our studies in mice have been confirmed in patients, demonstrating that obesity-induced changes can promote the expansion of preleukemic CHIP and lead to a severe MPN-like phenotype [15^{***}]. Given the shared pathogenetic mechanisms between obesity and CHIP, this review endeavors to investigate both preclinical and clinical evidence. Our focus is directed towards unraveling the intricate association between CHIP and obesity. Additionally, we delve into the consequential implications of this interaction on the pathophysiology of leukemia development.

OBESITY

Obesity, a prevalent global health concern, is associated with various conditions such as type 2 diabetes mellitus (T2DM), dyslipidemia, hypertension, insulin resistance, and cardiovascular diseases (CVDs)

[21,22]. The primary contributor to T2DM is obesity-induced insulin resistance, compounded by insufficient insulin secretion from pancreatic β -cells [21,22]. While adipose tissue serves as a crucial energy reservoir, its excessive accumulation fosters obesity by creating a pro-inflammatory, hyperlipidemic, and insulin-resistant milieu which is characterized by several metabolic abnormalities, including impaired glucose uptake by cells, elevated blood sugar levels, dyslipidemia (abnormal levels of fats in the blood), and often leads to the development of T2DM, obesity, and other metabolic disorders. This condition can increase the risk of CVD [21,23]. The BMI is a commonly utilized metric to assess overall body fat, determined by dividing an individual's weight in kilograms (kg) by the square of their height in meters. Typically, the conventional BMI range spans from 18.5 to 24.9 kg/m². Those with a BMI of 25 kg/m² or higher are classified as overweight, while individuals with a BMI of 30 kg/m² or higher are categorized as obese [24]. Additionally, to assess obesity comprehensively, factors like body fat distribution are crucial, prompting the utilization of tools such as WHR and computed tomography (CT) scanning [23]. Leptin, a peptide hormone exclusively secreted by adipocytes, plays a pivotal role in regulating body weight. Lack of leptin (Lep) or its receptor (Lepr) in both mice and humans leads to obesity, hyperglycemia, and insulin resistance [25]. In individuals with obesity, as well as in *Ob/Ob* mice, there is an increased accumulation of marrow adipocytes, forming fatty bone marrow (FBM). FBM negatively impacts bone mass by impeding osteoblast differentiation, thereby altering hematopoiesis [26,27]. Furthermore, bone marrow adipocytes secrete pro-inflammatory cytokines such as *IL-6*, *IL-1 β* , and *TNF- α* , contributing to the complex interplay between obesity and its associated health complications [28,29].

OBESITY, CHIP, AND LEUKEMIA-CURRENT EVIDENCE

CHIP emerges as a noteworthy risk factor for hematologic malignancies, yet the factors influencing clonal expansion and disease progression remain elusive [1,2]. Exploring the association between obesity and CHIP-associated diseases, in our recent study, we investigated individuals with CHIP-associated mutations. Analyzing a cohort of 47 466 participants from the UK Biobank, we found a correlation between CHIP mutations and a higher WHR, along with an elevated prevalence of coronary artery disease (CAD), hypertension, and hypercholesterolemia compared to those with lower WHR [15^{***}]. Notably, our research revealed a significant association between obesity, measured by both BMI

($n=46\,460$) and WHR ($n=47\,405$), and myeloid leukemia [15^{***}]. This observation aligns with recent studies demonstrating a compelling link between obesity and various types of leukemia, encompassing myeloid and lymphoid subtypes [13,14,16,30–35]. Numerous publications, including works by Larsson and Wolk [14], Kasim [35], and Castillo *et al.* [30], have consistently shown an elevated risk and incidence of AML and chronic myeloid leukemia (CML) in individuals classified as obese or overweight. Furthermore, MacInnis *et al.* [36] established an association between increased BMI and myeloid leukemia risk, emphasizing the multifaceted impact of adiposity.

A distinct subset of AML, acute promyelocytic leukemia (APL), also demonstrates a correlation with higher BMIs, leading to complications such as thrombohemorrhagic early death and increased treatment-related mortality. Importantly, pediatric AML patients with obesity exhibit inferior survival rates, as revealed in the CCG-2961 clinical trial. The intricate relationship extends beyond clinical outcomes, as individuals with *de novo* APL showcase elevated median BMIs and a higher incidence of obesity, reinforcing the potential link between obesity and myeloid leukemias [34,37–39]. In a recent investigation involving adolescents and young adults (AYAs) receiving treatment with Dana-Farber Cancer Institute (DFCI) consortium protocols for acute lymphocytic leukemia (ALL), high BMI was identified as being associated with increased toxicity, elevated nonrelapse mortality (NRM), and compromised overall survival [40^{*}]. Moreover, a separate study emphasized the pivotal role of inflammatory signals from FBM in promoting *DNMT3A*-driven clonal hematopoiesis, with the inhibition of this process achievable through the disruption of the IL-6 pathway [41^{***}]. Notably, obesity and an elevated BMI are recognized as risk factors for the development of multiple myeloma [42,43^{***}]. For instance, individuals with obesity face a 20% higher risk of multiple myeloma compared to their lean counterparts [44]. Collectively, these studies strongly indicate a potential correlation between obesity and the occurrence and fatality rates of various forms of leukemia. Notably, evidence points towards associations with AML, CML, chronic lymphocytic leukemia (CLL), ALL, multiple myeloma, and preleukemic conditions like myelodysplastic syndromes (MDS) [13,14,30, 45,46].

In light of the co-occurrence of CHIP-associated mutations in both myeloid and lymphoid malignancies, our findings align with existing literature showcasing a correlation between obesity and leukemia. These collective insights emphasize the

clinical significance of obesity in leukemia patient populations and underscore the need for further research to unravel the intricate mechanisms linking adiposity, clonal hematopoiesis, and hematologic malignancies.

Our experimental investigations in murine models validate the clinical observations within the patient cohort, establishing a proof of concept that alterations induced by obesity can instigate the expansion of preleukemic CHIP into a pronounced MPN-like phenotype. Utilizing several CHIP-associated genes, including *Tet2*, *Dnmt3a*, *Asx1*, and *Jak2*, our studies consistently demonstrated heightened clonal expansion and the manifestation of an MPN-like phenotype [15^{***}]. This universal effect across various mutation types suggests that obesity might universally catalyze or exacerbate CHIP-associated diseases (Fig. 1) [15^{***}]. Similarly, recent studies revealed that obesity induced by a high-fat diet (HFD) and the transplantation of bone marrow cells from mice lacking one copy of *Dnmt3a* (referred to as HET, for *Dnmt3a*^{+/-}, heterozygous) resulted in increased spleen size, leukemia development, and reduced lifespan in HET recipients on HFD compared to control groups [47^{***}]. Moreover, HET recipients on a HFD exhibit elevated levels of pro-inflammatory cytokines, including IL-6, both in plasma and within immune cells [47^{***}]. Concurrently, an elevation in pro-inflammatory cytokines such as IL-1 β , IL-6, and TNF- α , along with increased intracellular Ca²⁺ levels, was observed [15^{***}].

The intricate link between intracellular Ca²⁺ levels and adipocyte lipid metabolism are noteworthy. The regulation of lipid metabolism and triglyceride storage in adipocytes heavily relies on intracellular Ca²⁺ levels [48]. Elevated intracellular Ca²⁺ stimulates the expression of lipogenic genes while concurrently suppressing lipolysis, ultimately leading to increased adiposity [48]. We hypothesized that elevated intracellular Ca²⁺ levels might instigate aberrant signaling, potentially accelerating the onset of severe MPN. To explore therapeutic avenues, we treated *Ob/Ob* mice harboring *Tet2*^{-/-} preleukemic HSC/Ps with various agents, including metformin, pioglitazone, nifedipine, SKF-96365, MCC950, and anakinra, either individually or in combination [15^{***}]. Notably, the combination therapy outperformed individual treatments, indicating that a multipronged approach targeting multiple signaling pathways or environmental stimuli may be crucial for suppressing clonal hematopoiesis, particularly in the context of obesity. This combination therapy not only reduced the frequency of leukemic blasts, undifferentiated stem cells, HSCs, and granulocyte-macrophage progenitors but also mitigated arterial fatty deposits in *Ob/Ob* recipients

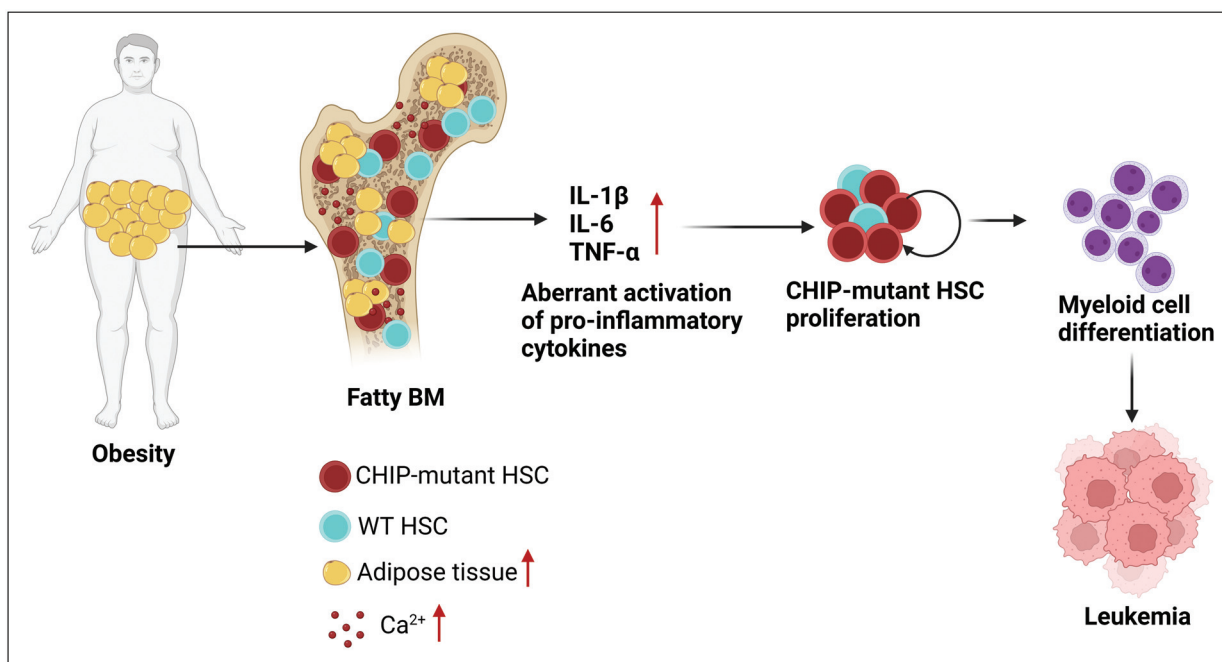


FIGURE 1. Illustration of the influence of obesity-induced inflammation on clonal hematopoiesis and leukemia development. In the obese bone marrow, inflammatory cytokines (IL-1 β , IL-6, and TNF- α) and heightened Ca^{2+} levels activate CHIP-mutant (HSCs). This activation leads to the proliferation of CHIP-mutant HSCs and their subsequent differentiation into myeloid cells, ultimately contributing to the development of leukemia. CHIP, clonal hematopoiesis of indeterminate potential; HSC, hematopoietic stem cells.

with CHIP, demonstrating the potential of inhibiting Ca^{2+} signaling in ameliorating CHIP-associated diseases [15²²]. Conducting follow-up studies involving the transplantation of specific myeloid populations in HFD-induced obese models [47²²], similar to our previous investigations utilizing *Tet2* cells on *Ob/Ob* mice [15²²], promises to be an intriguing avenue for further research.

Supported by clinical evidence, CHIP emerges as a high-risk factor for the development of hematologic malignancies, and its association with poorer prognosis extends even to individuals with solid tumors [23–29]. Our analysis of cancer genome atlas (TCGA) data further underscores the intricate connection between CHIP and obesity. Specifically, the mutation rate of CHIP-associated genes was markedly higher in patients with a BMI exceeding 30 kg/m² compared to those with a BMI of 25 kg/m² or lower across seven distinct cancer types, including breast (BRCA), lung (LUAD), ovarian, stomach (STAD), colon (COAD), low-grade glioma (LGG), and liver (LIHC) cancer. Concomitantly, obesity itself is implicated in an escalated risk across various tumor types. A large cohort of study reported that a substantial proportion (40%) of cancer diagnoses in the United States, were attributed to being overweight or obese [49]. Cancer stem cells in obese individuals may reprogram adipocytes (highly

accumulated in obese individuals BM) into cancer associated adipocytes, which subsequently secrete growth, inflammatory, fibrotic and angiogenic factors, and lymphokines [such as tumor necrosis factor (TNF- α), adiponectin, interleukin-6 (IL-6), IL-1 β , and vascular endothelial growth factor (VEGF)] that may promote tumor growth [50] or alter anticancer effects of chemotherapy [51]. IL-6, for instance, plays a role in sustaining proliferative signaling, mutation, tumor-promoting inflammation, invasion, metastasis, genome instability, and resistance to cell death [52²,53]. Similarly, IL-1 β is associated with invasion activation, metastasis, genome instability, mutation, tumor-promoting inflammation, and immune evasion [52²,54]. TNF- α is linked to proliferative signaling sustenance, evasion of growth suppressors, induction of angiogenesis, activation of invasion and metastasis, tumor-promoting inflammation, and immune evasion [52²,55]. VEGF stands out as the paramount regulator in vascular system development, and obesity significantly accelerates tumor growth through mechanisms entailing the heightened activation of VEGF pathways [56].

Crucially, these proinflammatory cytokines, which are elevated in both obese individuals and those with CHIP, suggest a potential synergistic effect between these conditions in promoting cancer

development [52[■],57–59]. The intersection of obesity and CHIP unveils a complex interplay that may significantly contribute to the pathophysiology of cancer, shedding light on the need for comprehensive investigations into these multifaceted relationships.

In summary, while emerging research has initiated the exploration of connections among obesity, CHIP, and the development of leukemia, the intricacies of the underlying pathophysiological mechanisms remain speculative and warrant further investigation. Insightful studies delving into inflammation within FBM have revealed intriguing overlaps in biomarker profiles with those observed in CHIP. These findings underscore the need for comprehensive investigations to unravel the intricate interplay between obesity, CHIP, and leukemia, aiming to elucidate the mechanistic links and potential therapeutic targets in this complex disease.

FUTURE PERSPECTIVES AND CONCLUSION

Obesity is associated with heightened risks of CVD, diabetes, insulin resistance, leukemia, and increased mortality compared to those with normal weight [1,2,21,22]. The connection between obesity and leukemia holds critical implications for prevention and treatment. Obesity emerges as a modifiable risk factor for leukemia, potentially offering a path for prevention. While current leukemia research often concentrates on expensive treatments, prioritizing prevention could prevent numerous cases. Robust studies are necessary to investigate whether obesity predisposes individuals to leukemia. Further research is essential to pinpoint the precise magnitude of the increased vulnerability to developing CVD, T2DM, leukemia, and ultimately, mortality in obese individuals with CHIP. This information is pivotal for deciding whether active CHIP screening is necessary in the obese population, considering the potential significant rise in risk. Adopting a healthier lifestyle, encompassing weight loss, balanced nutrition, and regular exercise, may diminish cancer risk. The exploration of pharmacological interventions, repurposing drugs, and contemplating bariatric surgery could provide valuable strategies for preventing leukemia and other obesity-related cancers.

In conclusion, epidemiological findings indicate an association between obesity and leukemia. Furthermore, individuals with leukemia who are obese tend to experience poorer outcomes and higher mortality rates. Conversely, the shared pro-inflammatory characteristics associated with both CHIP and obesity set off a chain of events, consequently heightening the risk of leukemia development (Fig. 1).

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Conflicts of interest

There are no conflicts of interest.

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