



Shared mechanisms of multimorbidity in COPD, atherosclerosis and type-2 diabetes: the neutrophil as a potential inflammatory target

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Multimorbidity is increasingly common in our ageing population. Drawing together research across chronic inflammatory disease is vital to enable effective medical intervention and prevent research becoming siloed <http://bit.ly/32l9hXs>

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ABSTRACT Multimorbidity is increasingly common and current healthcare strategies are not always aligned to treat this complex burden of disease. COPD, type-2 diabetes mellitus (T2D) and cardiovascular disease, especially atherosclerosis, occur more frequently together than expected, even when risk factors such as smoking, obesity, inactivity and poverty are considered. This supports the possibility of unifying mechanisms that contribute to the pathogenesis or progression of each condition. Neutrophilic inflammation is causally associated with COPD, and increasingly recognised in the pathogenesis of atherosclerosis and T2D, potentially forming an aetiological link between conditions. This link might reflect an overspill of inflammation from one affected organ into the systemic circulation, exposing all organs to an increased milieu of proinflammatory cytokines. Additionally, increasing evidence supports the involvement of other processes in chronic disease pathogenesis, such as cellular senescence or changes in cellular phenotypes.

This review explores the current scientific evidence for inflammation, cellular ageing and cellular processes, such as reactive oxygen species production and phenotypic changes in the pathogenesis of COPD, T2D and atherosclerosis; highlighting common mechanisms shared across these diseases. We identify emerging therapeutic approaches that target these areas, but also where more work is still required to improve our understanding of the underlying cellular biology in a multimorbid disease setting.

Introduction

Multimorbidity is defined as the presence of two or more chronic conditions in the same person. It is increasingly common in our ageing population, affecting nearly 40% of European citizens [1] and is extremely costly in terms of healthcare resource use [2]. Some conditions co-occur more commonly than others and COPD is often present in multimorbid disease clusters with cardiovascular disease (CVD), especially atherosclerosis and type-2 diabetes mellitus (T2D) [3, 4], conditions associated with an immense

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global burden of ill health. These conditions share multiple risk factors, including age, smoking, obesity, physical inactivity and poverty [5, 6]. For example, smoking cigarettes is clearly associated with COPD pathogenesis, but also shares a robust association with diabetes and atherosclerosis after multivariable adjustment [7, 8]. However, smoking is neither necessary nor sufficient to cause any of these diseases and even after accounting for overlapping risk factors, the incidence of COPD, T2D and atherosclerosis presenting in the same person is still higher than expected [3, 4, 9].

There is great interest in understanding why multimorbidities cluster in individuals [10]. Network analyses have identified multiple shared molecular pathways and genetic associations between COPD, cardiovascular diseases and T2D [11, 12], perhaps alluding to common mechanisms. When present, these mechanisms might lower susceptibility to cause or increase the progression rate of components of the clustered diseases. This provides the opportunity for shared therapies that cut across traditional disease silos; a holistic approach to pathology permitting a holistic approach to patient care.

Multimorbidity is more common with increasing age and is associated with inflammation and cellular dysfunction. Targeting these processes (cellular ageing, inflammation and cellular dysfunction) may offer novel treatments across multimorbidities or allow us to utilise the drugs we already have for new indications (“repurposed” drugs). For example, statins, used to lower cholesterol as part of a CVD risk-reduction strategy are under investigation across a wide range of disease areas, including infections [13], COPD [14] and dementia [15]. However, research into shared mechanisms of multimorbidity are limited as most clinical observational studies and interventional trials focus on a single disease entity.

This review discusses known and emerging mechanisms of disease pathogenesis, which could be shared across multimorbidity, using COPD, atherosclerosis and T2D as exemplar diseases. While these are among the most common noncommunicable chronic diseases globally, they also present as risk factors for each other and there is an increased prevalence of COPD, atherosclerosis or T2D in patients with a diagnosis of any of these conditions [16–18]. In particular, this review will focus on the shared role of the neutrophil in these diseases. While neutrophils are not the only cells that play a role in COPD, atherosclerosis and T2D, they are the most abundant immune cells found in lung secretions of patients diagnosed with COPD and their products have been shown to cause all facets of lung disease in animal and cell models (from chronic bronchitis to emphysema); correlating with disease severity and decline [19, 20]. Recent advances in imaging technologies have also identified that neutrophils are present in the lipid core of human plaques and neutrophilia correlates with the number of rupture-prone lesions [21] and, although less studied, diabetic patients, rats and mice clearly demonstrate consistent defects of neutrophil chemotactic, phagocytic and microbicidal activities [22]. Together, this suggests neutrophils may provide a link between these three disease areas.

An overview of mechanisms for multimorbidity

There are multiple mechanisms that play a role in the development of COPD, atherosclerosis and T2D including: altered function of cells such as endothelial cell and neutrophils [23–25]; mitochondrial defects [26]; metabolic changes [27]; increased age [25, 28]; and increased inflammation [29, 30]. Despite overlapping mechanisms and risk factors, targeting a known manifestation of one disease does not always improve outcomes in another. For example, targeting glucose levels in patients with T2D [31] or improving airflow obstruction in COPD [32] does not necessarily improve cardiovascular mortality. There is an urgent need to further our understanding of the potential links between these disease areas to guide therapies for multimorbid patients. This review will focus on three main, yet overlapping, putative common mechanisms observed in COPD, atherosclerosis and T2D, while appreciating other mechanisms do exist as detailed above.

First, COPD, atherosclerosis and T2D are more common in an ageing host and “accelerated ageing” has been proposed as a pathogenic mechanism in these diseases, including ageing of individual cells resulting in a complex phenotype termed “senescence” (figure 1a) [33]. Cellular senescence was initially described as a process whereby cells entered permanent cell cycle arrest, allowing aged cells to be cleared. However, our understating of senescence has expanded, especially regarding nonreplicating cells, such as neutrophils (figure 1a) [34]. It is now recognised that senescent cells are diverse, and their function and fate may depend on the strength and duration of the stressor they are exposed to, including oxidative stress and DNA damage. These may lead to cell repair, immune clearance of the senescent cell or phenotypic changes in senescent cells that endure in tissues (as reviewed by VAN DEURSEN [35]).

Secondly, and linked to theme 1, atherosclerosis, COPD and T2D are associated with chronic systemic inflammation (figure 1b). Inflammation is both caused by and drives many chronic noncommunicable diseases, including COPD, atherosclerosis and T2D. Increased systemic inflammation has been reported in multimorbid patients when compared to those with a single disease or age-matched healthy volunteers [36, 37]. It is important to note that while similar patterns of inflammation, including neutrophilic

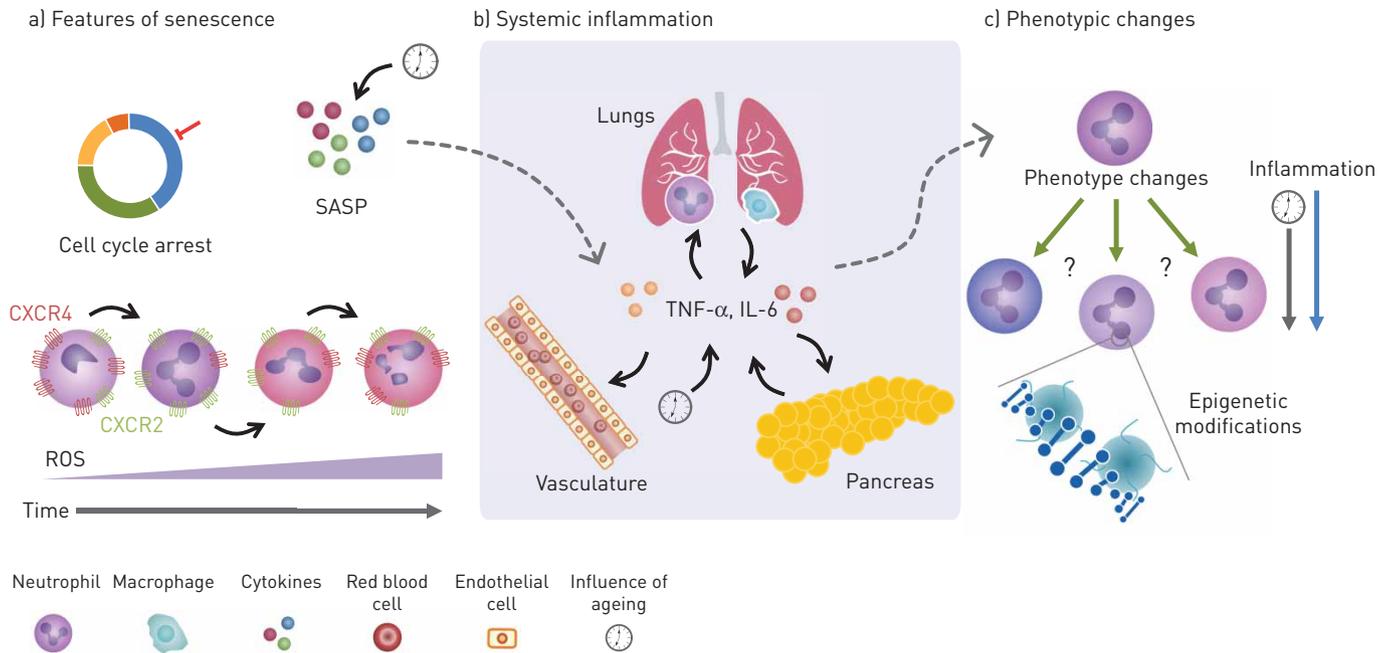


FIGURE 1 Potential common inflammatory mechanisms between COPD, type-2 diabetes mellitus (T2D) and cardiovascular disease (CVD). a) Senescence is described as changes that result in cell-cycle arrest and is associated with a senescence-associated secretory phenotype (SASP). The SASP leads to increases in inflammatory cytokines both locally and systemically (dotted arrow). This process in neutrophils, while slightly different to typical senescence, might be characterised by increases in chemokine receptor CXCR4 and decreases in CXCR2. This may change the way neutrophils behave and contribute to the damage seen in these diseases. These changes are also associated with increased reactive oxygen species (ROS) production, causing an oxidant imbalance and increasing damage. b) Inflammatory cytokines released by lung epithelial cells and immune cells can enter circulation from the lung in COPD. Likewise, these cytokines can also be released by immune cells in the pancreas in T2D and from endothelium and plaques in CVD. Increases of these cytokines also occur with age and can influence the cytokine milieu, leading to potential changes in neutrophil phenotype (dotted arrow). c) Changes in neutrophil subpopulations (or cell phenotype) caused by inflammation may also play a role in the crosstalk between COPD, CVD and T2D by contributing to tissue damage. There is also some evidence to suggest epigenetic changes may perpetuate the disease state causing long-term maladaptive changes to cellular phenotype and function, influenced by both inflammation and ageing. TNF: tumour necrosis factor; IL: interleukin.

inflammation, are observed in COPD, atherosclerosis and T2D [38–40], inflammation is a mechanism that plays a role in numerous chronic and acute diseases. Systemic crosstalk of inflammatory cytokines might occur through “overspill”, where inflammatory proteins produced in one organ can spread system-wide (figure 1b), impacting on the development or progression of other inflammatory diseases. The proximity of atherosclerotic plaques to the bloodstream provides obvious routes for systemic inflammation, but this mechanism has also been described in COPD and T2D [39, 41].

Thirdly, immune cell activation and dysfunction have been described in COPD, atherosclerosis and T2D, and linked with adverse disease outcomes (figure 1c). Clearly, neutrophils are not the only cell that show altered functionality in these three diseases, but the fact that neutrophil dysfunction occurs in COPD, atherosclerosis and T2D may provide novel links between these diseases [22, 42, 43]. This cellular dysfunction may represent a physiological response to a pathological event, but there is increasing evidence that chronic inflammation (as seen in COPD, atherosclerosis and T2D) can contribute to a shift in immune cell phenotype or subpopulation, changing the cell’s usual response to an inflammatory insult, and increasing the potential for damage (figure 1c). This may or may not reflect cellular senescence, but epigenetic modification may be one potential mechanism for this long-term phenotypic shift [44], which could impact on multimorbidity.

These themes will be explored in more detail below. The overlapping nature of these proposed mechanisms have been summarised in figure 1.

Theme 1: accelerated ageing in multimorbidity

Ageing-associated cellular senescence is also seen in the immune system, termed “immunosenescence”, with less effective pathogen containment and killing, enhanced inflammation and inefficient repair processes [25, 45].

The parallels between age and chronic disease are manifold, leading to the hypothesis that chronic conditions reflect a state of advanced or accelerated ageing. Both age and chronic disease are associated

with primary hallmarks of ageing-related dysfunction: genetic instability, epigenetic alterations, loss of proteostasis and telomere shortening [28].

DNA instability has long been noted in COPD [46] and more recently described in T2D [47] and atherosclerosis [48]. Epigenetic modifications, including changes in histone deacetylase 2 activity, have been identified in all three conditions [49–51]. A loss of proteostasis and autophagy has been associated with emphysema in COPD [52] and also contribute to T2D [53] and atherosclerosis progression [54]. Studies have described a reduction in relative telomere length (RTL) in leukocytes in COPD, atherosclerosis and T2D, but this has not been universally replicated [55, 56]. A reduction in RTL was observed in peripheral lymphocytes in patients with COPD [57] and in lung epithelial cells exposed to cigarette smoke *in vitro* [58]. However, no significant difference in RTL was found in lung biopsies from patients with COPD compared to healthy controls [59], suggesting that these changes might be cell-type specific. Similarly, in T2D and atherosclerosis, RTL has been shown to be reduced in circulating leukocytes but with variable penetration across disease severity [60, 61].

These changes are thought to be due to the time-dependent accumulation of cellular damage; however, it remains unclear whether these changes occur before the onset of age-associated disease or as a result of disease, in all or only a subset of cells, and to what extent these changes are reversible.

Multimorbidity is associated with an increased burden of senescent cells, compared to health [62]. Senescent cells can be identified by senescence-associated β -galactosidase (SA β G) activity [63] (although not all senescent cells express SA β G [64]) or by the cocktail of inflammatory cytokines, growth hormones and proteinases they secrete, called the senescence-associated secretory phenotype (SASP) [65].

An increased presence of senescent cells, SA β G activity and the SASP have been described in atherosclerosis by both vascular smooth muscle cells and endothelial cells around atherosclerotic plaques *in vitro* and *in vivo* [66, 67]. In T2D, a high glucose level has been shown to drive premature senescence *in vitro* in endothelial cells, renal mesangial cells, adipose-derived stem cells and fibroblasts [68]. In COPD, a higher burden of senescent cells has been described in pulmonary vessels, especially those with vascular remodelling that has been associated with pulmonary hypertension, suggesting an interesting link between COPD and vascular disease [69]. A similar increase in senescence of the smooth muscle that surrounds pulmonary arteries in these patients was also observed, providing further links between vascular senescence, atherosclerosis and COPD [69]. Small sections of regulatory RNA, known as microRNAs (miRs) have been implicated in promoting cellular senescence and miR-34a has been associated with fibroblast senescence in COPD [70] atherosclerosis [71] and T2D [72], demonstrating a common link between diseases.

Circulating cells, such as neutrophils, may convey pathologies and enhance ageing processes between tissues, providing a link between multimorbid conditions. Moreover, there is evidence to support reverse transmigration of neutrophils from tissue into the blood, suggesting that any local senescent phenotype could be conferred systemically by neutrophils to other organs [73]. Neutrophil senescence in humans leads to distinct functional changes, including migration, phagocytosis and NETosis, which impair the host response to infection, while simultaneously increasing the potential for host damage through inflammation, reactive oxygen species (ROS) and proteases activity [74, 75]. Human neutrophil migratory accuracy reduces with age [25], a feature that is exaggerated in COPD [76] and potentially increases tissue damage through migration-associated release of proteases and ROS [25, 77]. Reduced neutrophil migratory accuracy and reduced bacterial killing has been identified in hyperglycaemia and replicated *in vitro* following neutrophil exposure to serum from diabetic patients [78]. *In vitro*, neutrophils have also been implicated in cell-to-cell crosstalk, with neutrophil microvesicles associated with inducing classical pathological changes in diabetic cardiovascular pathology [79] and COPD [80]. Furthermore, age-related changes seen in the muscle (*e.g.* sarcopenia) are also associated with an increase in neutrophil numbers: one study found a correlation between frailty and increased baseline neutrophil numbers [81] and a potential decline in neutrophil function [82], providing evidence for the role of neutrophil dysfunction to cause systemic ageing pathologies.

There is evidence that removal of senescent cells (senolysis) has a positive effect on age-related diseases [83] and age-related physical dysfunction in mice [84], including sarcopenia, cataracts and atherosclerosis *via* an inducible transgene that can deplete senescent cells [85, 86]. These transgenic approaches do not directly translate to a viable treatment in humans but targeting other features of senescence (such as SA β G activity) may be a therapeutic option. A first-in-human study has suggested that senolytics are safe and improved physical function in patients with idiopathic pulmonary fibrosis [87]. Preclinical studies in T2D identified that replicative-senescent β -cells actually contributed more to insulin production than nonsenescent cells [88], yet targeting senescent cells intermittently in mice still improved glucose tolerance and increased insulin sensitivity [89]. However, not all results are positive; in atherosclerosis, quercetin (a senolytic) was associated with nonsenescent cell death in primary human coronary artery endothelial cells

[90], despite earlier studies suggesting potential benefits [91]. This suggests caution is required within this field and more trials are eagerly awaited.

Theme 2: systemic inflammation and overspill in multimorbidity

Driven by the primary hallmarks of ageing, there is low-grade inflammation in older individuals which is similar to the systemic inflammation seen in chronic disease, a phenomenon termed “inflammaging” [29]. In disease, this may be further exaggerated and although many inflammatory cytokines have been shown to be raised in COPD, atherosclerosis and T2D, the significant overlap present in inflammation means it is difficult to identify an initiating signal. Tumour necrosis factor (TNF)- α increases with increasing age [30] and has been repeatedly implicated in COPD, atherosclerosis and T2D (figure 2) and, therefore, will be used as example mediator of systemic inflammation; however, other cytokines, including interleukin (IL)-6 and IL-1 β , have multiple systemic effects and are likely to be involved in multimorbidity as well (reviewed

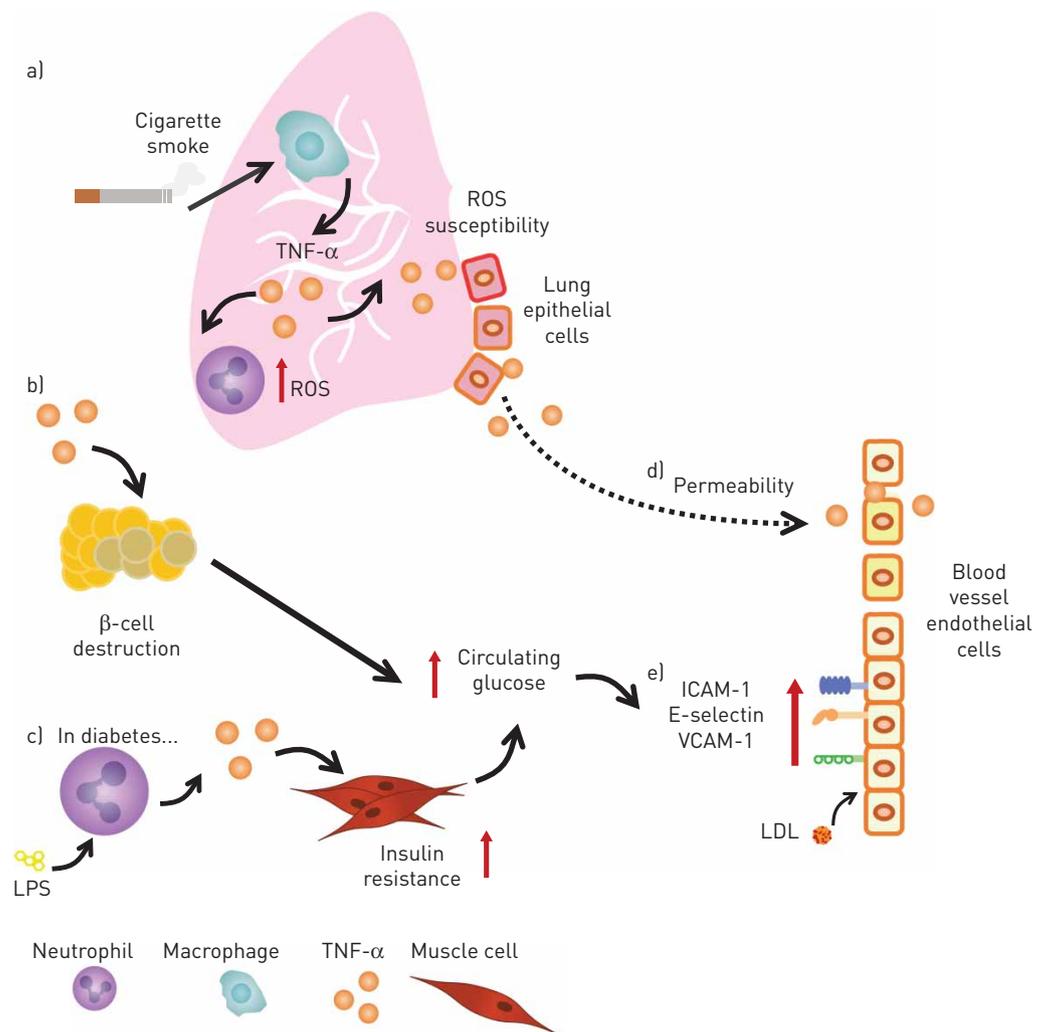


FIGURE 2 A summary of the multiple influences of tumour necrosis factor (TNF)- α across COPD, type-2 diabetes mellitus (T2D) and atherosclerosis. TNF- α is produced by multiple cell types in various compartments of the body. a) Cigarette smoke increases TNF- α production by alveolar macrophages, leading to enhanced reactive oxygen species (ROS) production by neutrophils and increased ROS susceptibility of lung epithelial cells. b) From the perspective of T2D, circulating TNF- α can cause β -cell destruction that in turn leads to increases in blood glucose levels. c) Enhanced TNF- α production by neutrophils exposed to lipopolysaccharide (LPS) in patients with diabetes promotes insulin resistance, which also increases blood glucose levels. Both the mechanisms in b and c lead to glucose-driven adhesion molecule expression in the vasculature. d) TNF- α also increases vascular permeability and in COPD this can be from either TNF- α produced in circulation or diffusion of pulmonary TNF- α . e) In the vasculature, TNF- α increases adhesion molecules, such as intracellular adhesion molecule (ICAM)-1, E-selectin and vascular adhesion molecule (VCAM)-1, increasing the likelihood of immune cell infiltration. It also causes the promotion of low-density lipoprotein (LDL) uptake by endothelial cells.

elsewhere [30]). The inflammasome (a multiprotein complex that regulates inflammatory cytokine production, including IL-1 β) has also been described as being of importance in some models of COPD [92], atherosclerosis and T2D, and is again reviewed elsewhere [93].

Some studies have demonstrated an association with COPD severity and concentrations of systemic TNF- α , for example SINGH *et al.* [38], although this has not been universally replicated [94], potentially indicating heterogeneity within patient populations. Macrophages produce TNF- α in response to cigarette smoke extract *in vitro* [95] and *in vivo* using smoke-exposed rats (figure 2a) [96]. Alveolar macrophages appear to require exposure to both a Toll-like receptor agonist and cigarette smoke extract *in vitro* to increase TNF- α production [97], a mechanism relevant in the lungs, where both smoke and bacteria are present [98]. While TNF- α is also present in smokers without COPD, multiple studies investigating inflammation in COPD and smoking do not account for the fact that only a proportion of smokers develop COPD. Despite this, TNF- α is associated with multiple facets of COPD in murine and cell models, including: mucus hypersecretion [99]; matrix remodelling [100]; repair and maintenance contributing to emphysema [100]; and corticosteroid insensitivity through the reduction of histone deacetylase-2 activity [101]. While these data suggest that TNF- α has a role in the pathogenesis of COPD, potentially exaggerated in patients with multiple lung infections, much of this evidence comes from animal models.

TNF- α concentrations are also elevated in the plasma of patients with an acute myocardial infarction [102] and in the adipose tissue and serum of obese patients with CVD [103]. TNF- α is implicated in insulin resistance and β -islet cell dysfunction in murine models, suggesting that IL-1 β and TNF- α treatment result in a loss of the transcription factor FoxO1, which is required for maintaining β -cell identity (figure 2b) [104, 105]. TNF- α murine knockout models are more resistant to developing atherosclerosis [106], emphysema [107], endothelial dysfunction in diabetes [24] and late-stage diabetic retinopathy [108]. Neutrophils isolated from patients with T2D release larger amounts of proinflammatory cytokines, including TNF- α and IL-6 at baseline and in response to lipopolysaccharide (LPS) stimulation (figure 2c) [39]. It is possible that this response is similar to the enhanced TNF- α production in response to LPS seen in alveolar macrophages from patients with COPD [98], but further work is required to fully understand the impact of this neutrophil response to disease pathology. Furthermore, neutrophils isolated from patients with COPD released more proinflammatory cytokines in response to LPS in the presence of TNF- α [109], demonstrating a possible feedback loop between raised TNF- α production during infection and further proinflammatory signalling by neutrophils in COPD and T2D.

Collectively, these data suggest that TNF- α is elevated in COPD, atherosclerosis and T2D, and may contribute to individual diseases. One major caveat with many of the discussed studies is the lack of stratification of patients based on their comorbidities, and therefore it is likely that patients in these studies may already have a combination of COPD, atherosclerosis or T2D. Without this detailed information, it is impossible to fully understand the mechanisms underpinning pathology of a single disease, and to dissect the potential shared mechanisms in multimorbid patients. However, if inflammatory cytokines are to have a major role in multimorbidity, inflammation in one compartment needs to influence other compartments in the body. "Overspill" may be a means to achieve this, potentially unifying disease-related processes, and again although other cytokines are implicated in this, TNF- α is used as an example.

Evidence for overspill

Overspill is readily envisaged in atherosclerosis, where vascular plaques permit cytokines to enter the bloodstream causing systemic inflammation [36]. However, overspill is also highly likely during local tissue inflammation (*e.g.* in the lung) with increased vascular permeability permitting the spread of inflammation into the systemic circulation (figure 2d) [110, 111]. Increased circulating inflammatory cytokines also correlate with frailty and the development of sarcopenia in several epidemiological studies [82], providing a basis for chronically raised cytokine levels to influence multiple areas of the body.

In patients with COPD, the levels of IL-1 β and TNF- α in the sputum of patients with COPD negatively correlated with body mass index, indicating a potential link between local disease in the lungs and systemic low-grade inflammation in the form of obesity [112]. However, in this study, sputum TNF- α concentrations did not correlate with plasma level [112], but this may reflect the short half-life of free TNF- α in biological fluids and the dilution of biological fluids, as seen in induced sputum and lavage samples [113], which are important considerations when interpreting systemic cytokine levels in disease. Despite the dilution of any cytokine overspill from the lungs into the blood, some studies suggest that systemic inflammation could drive the pathologies of COPD: a study of never-smoking patients with rheumatoid arthritis described some participants as having clear evidence of emphysema on computed tomography scans [114]. The exact mechanism of this observation has not been investigated but is likely to involve a combination of genetic and environmental factors alongside systemic inflammation. A study

using a mouse model of COPD further supported the ability of lung inflammation to influence the vascular environment, as mice overexpressing IL-18, another inflammatory cytokine, developed impaired glucose tolerance over time [115].

A mechanism that could unify changes across diseases is the ability of TNF- α to increase the expression of capture receptors and adhesion molecules on the surface of blood vascular endothelial cells that are necessary to support leukocyte ingress into inflamed tissue [116]; a response that can be blocked using TNF- α inhibitors (figure 2e) [117]. Systemic changes in the expression of adhesion molecules by blood vascular endothelium also occur in patients with T2D, potentially preceding the onset of T2D [118] and modification of vascular adhesion molecule-1 has been linked to early atherosclerosis progression in cell lines from *in vitro* and *in vivo* studies [119, 120]. TNF- α also increased low-density lipoprotein uptake in vascular endothelium *in vitro*, a process known to promote atherosclerosis and further proinflammatory signalling (figure 2e) [40]. In patients with COPD, the influx of leukocytes into the lungs contributes to lung damage through protease and ROS, a process which TNF- α can enhance, further contributing to damage [121]. Furthermore, TNF- α increased the susceptibility of pulmonary vascular endothelium to ROS *in vitro* by reducing glutathione levels intracellularly, presenting a mechanism of enhanced damage of pulmonary endothelium in oxidative environments (figure 2a) [122].

Matrix remodelling promoted by TNF- α through matrix metalloproteinases (MMPs) may also provide a link between COPD, CVD and T2D, albeit through different mechanisms. Matrix remodelling through increased MMP-9 and MMP-2 activity have been linked to increased lung destruction and inflammation in COPD [100]. In CVD, MMPs have been shown to promote the occlusion of blood vessels during atherogenesis by enhancing blood clotting [123] and MMP-9 specifically has been linked with familial history of T2D and with increasing insulin resistance [124]. Matrix remodelling has also been associated with ageing in multiple studies, potentially due to age-related loss of functionality of cells responsible for maintaining the extracellular matrix [125].

These data present complex but unifying mechanisms of systemic inflammation in the pathogenesis of COPD, T2D and atherosclerosis. However, data from clinical studies of inflammatory processes have ambivalent results. Retrospective studies have identified a potential benefit for patients with T2D receiving anti-TNF- α therapy for rheumatoid arthritis [126], although no data currently exist to support the use of anti-TNF therapy for the sole purpose of treating T2D. Anti-TNF- α therapy (again used for rheumatoid arthritis) has also been reported to reduce hospitalisation due to COPD exacerbations [126], but a trial of specific anti-TNF- α therapy (infliximab) in unselected patients with COPD was not associated with improvements in health or lung function [127]. Patients on anti-TNF- α therapy for psoriasis were noted to have a reduced incidence of myocardial events in population studies [128] and patients with rheumatoid arthritis with similar treatment showed a reduction in aortic stiffness, linked to atherosclerosis progression [129]. Although etanercept, another TNF- α inhibitor, reduced systemic inflammation following an acute myocardial infarction, increased platelet activation was noted (a key feature of atherosclerosis) with no change in peripheral vasomotor or fibrinolytic function [130].

Trials of other interventions to reduce the activity of specific inflammatory cytokines have also been equivocal. Patients with a previous myocardial infarction treated with an anti-IL-1 β medication, canakinumab, demonstrated a reduction in cardiovascular events over the study period, but with no reduction in all-cause mortality [131], and the same drug provided no clinical benefit in COPD [132]. Targeting the IL-1 β receptor has also proved inefficacious in COPD, with the most common adverse event being worsening of COPD symptoms [133]. As with other re-purposed interventions, the timing, patient group, dose and duration of treatment may be fundamental to determining a response in multimorbidity and it might be that selected populations would benefit through offering a personalised medicine strategy, if they could be identified. Another important consideration is the bioavailability of TNF- α , or indeed other inflammatory cytokines, in the circulation. This could be influenced by free TNF- α (usually referred to as soluble TNF- α), membrane-bound or vesicular TNF- α , with each potentially influencing potency and the ability to be inhibited in the circulation [134].

Theme 3: altered cellular subpopulations and processes in multimorbidity – the lung as a central hub

Immune cell subtypes have been recognised for many years and have been implicated in chronic disease and multimorbidity. Initially, neutrophils were excluded from this concept, considered too short lived and too restricted in their responses to impact across disease clusters. However, emerging evidence has questioned this, and proposed a novel hypothesis to explain why neutrophils and damaged lungs may place the host at a particularly increased risk of other chronic inflammatory illnesses [73]. While other cell types are involved in the development of multimorbidity, the potential role of the neutrophil and the lungs will be explored here.

Neutrophils are known to exist in three states; quiescent, primed and activated, and in order to release proteases or build the reduced nicotinamide adenine dinucleotide phosphate oxidase (NOX) complex needed to generate ROS [135], neutrophils require significant stimulation to move from the quiescent to the activated state [136]. This system exists to prevent unwarranted host damage. Once cells become primed, activation occurs more readily and for several years it has been recognised that neutrophils can oscillate between these states, depending on the environment. Recent, elegant *in vivo* studies have shown that in health, human neutrophils do not coalesce in the lungs (as previously suggested) and instead pass through the pulmonary capillaries only marginally slower than red blood cells [137]. However, the transit time is much slower if neutrophils are primed, as has been shown in patients that have inflammatory lung diseases such as COPD [138] or even low-grade inflammation [139]. The reason for this slower transit time is thought to reflect that primed neutrophils adopt cytoskeletal changes which makes them stiffer and less deformable compared to quiescent cells, impacting on their ability to squeeze through the narrow capillaries. However, when neutrophils are artificially primed *ex vivo* and then reintroduced to the circulation, this retention signal in the lungs is steadily lost, suggesting these cells are either cleared from the body or are deprimed [137]. Forced mechanical deformation *ex vivo* of neutrophils has been shown to deprime neutrophils [140], and it has been suggested that the lungs may form an important site where neutrophils, primed through exposure to circulating inflammation, may return to the quiescent state because of mechanical modulation in the tight pulmonary capillaries. The narrow pulmonary vasculature might, therefore, have important functions for immunomodulation.

In chronic lung disease, the slower transit time persists, suggesting primed cells remain in the systemic circulation, perhaps caused by the pulmonary vasculature destruction and remodelling present in COPD (preventing depriving by mechanical manipulation of neutrophils) and the proinflammatory environment of the lungs [138]. These primed cells would be much more susceptible to respond to an inflammatory signal in another organ, perhaps linking lung disease with other chronic inflammatory pathologies. Furthermore, human neutrophils can reverse migrate *in vitro* through the endothelium and back into the circulation [73], after sampling inflamed tissues. These previously tissue-bound cells maintain a proinflammatory and apoptotic-resistant phenotype, which could provide a link between chronic inflammation and systemic damage in COPD, atherosclerosis and T2D as they circulate around the body [73]. This mechanism, however, would rely on COPD or associated lung damage being the initiator disease. While hypothetically low-grade inflammation may prime neutrophils, and subsequently add to the burden of disease and contribute to the pathogenesis of multimorbidity [139], this field is in its infancy and the evidence (although generally supportive) remains patchy.

There is increasing evidence for neutrophil heterogeneity and plasticity in terms of cellular responses and surface expression of receptors, with both pro and anti-inflammatory subpopulations described (figure 3) [141, 142]. An anti-inflammatory and pro-resolving neutrophil phenotype have been described, where the secretion of α -defensins modifies the inflammatory response of macrophages [143] and neutrophils can also modify angiogenic processes with the secretion of MMP-9 [144], as well as their typical proinflammatory responses. It is unclear whether the presence of chronic inflammatory disease in an ageing host leads to a loss of neutrophil plasticity, a change in phenotype or merely a priming of cells, but there is clear evidence of increased neutrophil activity and dysfunction across COPD, atherosclerosis and T2D (as described earlier) and studies are describing differences in neutrophil populations between patients with COPD and controls [145] and in murine models of atherosclerosis [146], although this field is in its infancy.

Key cellular processes may also link multimorbidities. ROS are known to cause damage in biological systems, including within settings of chronic inflammation [147], and are linked to ageing as described by the free radical theory of ageing [148]. Neutrophils have enhanced ROS responses in patients with COPD, especially with bacterially induced exacerbations [149, 150]. Raised markers of oxidative stress have been identified in the sputum and plasma of patients with COPD, related to increased ROS and also decreased antioxidant activity [151]. This increase in ROS production in the lungs presents a potential mechanism for local damage in patients with COPD. There is also evidence of systemic ROS activity in COPD, potentially predisposing to events seen in the pathogenesis of atherosclerosis and T2D [151, 152]. In atherosclerosis, upregulation of NOX2 expression has been described before atherosclerosis develops [153] and in diabetes, ROS production can contribute to β -cell destruction, as these cells are particularly sensitive to ROS-induced apoptosis [154]. NOX1 has also been shown to play a role in diabetes-associated complications from human *in vitro* and murine *in vivo* experiments: human aortic endothelial cells and cells in the renal cortex upregulate NOX1 levels in response to hyperglycaemic conditions [155] and NOX1 inhibition *in vivo* is associated with a reduction in atherosclerosis and reduced adhesion of inflammatory cells to the vascular wall [156]. Data sourced from the National Institutes of Health (ClinicalTrials.gov) show several trials of antioxidants in COPD,

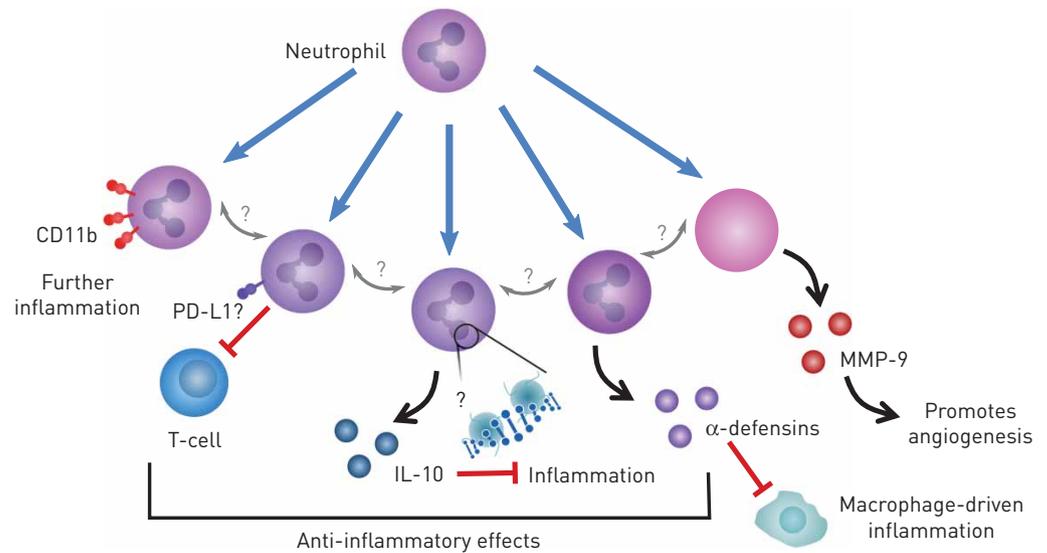


FIGURE 3 Neutrophil phenotypes can be altered, and these changes may impact the pathogenesis of other diseases. Several neutrophil phenotypes are described showing the plasticity of the neutrophil. Increasing surface expression of CD11b has been linked with further inflammation and therefore a proinflammatory phenotype. Several anti-inflammatory phenotypes have also been described: the suppression of T-cell function; the release of an anti-inflammatory cytokine interleukin (IL)-10; and the release of α -defensins shown to inhibit macrophage-driven inflammation by preventing mRNA translation. Neutrophils also release matrix metalloproteinase (MMP)-9, which contributes to matrix remodelling and promotes angiogenesis impacting on vascular remodelling in disease. This shows a broad array of inflammatory functions by the neutrophil. PD-L1: programmed-death ligand 1.

diabetes and atherosclerosis, suggesting there remains great interest in mitigating ROS-related damage in chronic disease.

Interwoven processes require an integrated approach

Current management strategies for COPD, T2D and atherosclerosis have improved outcomes for patients but have not halted disease processes. This might suggest we need a new therapeutic approach. The mechanistic themes outlined above (ageing and senescence, inflammation and cellular dysfunction) appear interwoven; each associated with each other. There are clear links with altered cellular functions, inflammation and senescence, with impaired immune cell surveillance associated with a higher burden of senescent cells [157], a greater burden of SASP and the associated systemic and tissue inflammation [36, 41, 112]. Once present, inflammation, including $\text{TNF-}\alpha$, serves as an important priming agent for the oxidative burst [121]. ROS can damage cells and tissues, instigate further proinflammatory responses and exacerbate DNA damage. This proinflammatory milieu is associated with epigenetic modification and RTL shortening, a loss of proteostasis and a greater burden of senescent cells, in what is considered a continuing cycle of damage and disease progression. These processes are more apparent in our globally ageing population. However, more research is needed. Not all environmentally exposed individuals develop disease (for example, not all smokers develop COPD or atherosclerosis), although there are clear clusters, not all chronic inflammatory conditions appear together, and age or environmental exposures alone do not predict those at most risk [10, 31]. It is unclear whether the biological hallmarks of ageing all present together in susceptible individuals, or whether one hallmark precedes the others. It is also unclear whether biological ageing is ubiquitous across tissues, or whether certain tissues, such as the lung with its high exposure to environmental toxins and bacteria, are more affected. The cellular dysfunction seen in COPD, atherosclerosis and T2D share common features, including altered neutrophil migratory dynamics and impaired phagocytic responses, but do not seem to only reflect a senescent cell. Perhaps it is the combination of chronic pulmonary and systemic inflammation, the presence of hyperglycaemia and the inability to deprime cells in damaged lungs in an ageing host that makes the combination of these conditions so dangerous for an individual.

A challenge for treating multimorbidity also lies in healthcare procedures, as the initial disease diagnosis usually dictates the treatment pathway. If these interwoven inflammatory mechanisms underly the disease process, identification of the predominant pathway, as opposed to the first diagnosed disease, may improve treatment outcomes. This is also met with a relatively poor understanding of early disease with increasing

TABLE 1 Key ongoing or recently completed interventional trials in COPD, type-2 diabetes mellitus (T2D) and cardiovascular disease (CVD) related to the mechanisms of ageing, inflammation and cellular phenotype

Study name	Study identifier	Study overview	Disease area	Phase
Ageing				
HIV-related Accelerated Aging of the Airway Epithelium	NCT01974219	Observational study investigating the role HIV may play in the accelerated ageing of the small airway endothelium and development of COPD	COPD	N/A
Pulmonary Rehabilitation Innovation and Microbiota in Exacerbations of COPD (PRIME)	NCT03701945	An interventional study to investigate how the microbiota of the lungs contributes to pulmonary ageing and AECOPD, and the impact pulmonary rehabilitation has on AECOPD	COPD	N/A
In Utero Smoking and Premature Cellular Senescence	NCT01865435	A trial to investigate telomere length as a surrogate for cellular senescence in peripheral lymphocytes obtained from the cord blood of newborns from mothers with absence of smoking or smoking >5 cigarettes per day	COPD	N/A
Nutritional and Functional Changes in Heart Failure and COPD	NCT01787682	An interventional trial to investigate gut absorption in patients with CVD or COPD and if this can be improved with a high protein supplement to reduce muscle loss in these patients	CVD, COPD	N/A
Cardio-vascular Protective Effects of Wolfberry in Middle-aged and Older Adults	NCT03535844	A single-blind 16-week study investigating the protective effects of Wolfberry in positively changing endothelial function and lipidomic profiles	CVD	N/A
Regulation of Endothelial Progenitor Cells by Short-Term Exercise (EPC-Ex)	NCT01169831	An open interventional study investigating the effects of exercise on endothelial progenitor cell numbers in sedentary older adults and older endurance athletes	CVD	N/A
Prevention of Cardiovascular Stiffening With Aging and Hypertensive Heart Disease (LVH)	NCT03476785	An open interventional trial to assess how exercise intervention prevents the age-related stiffening of the left ventricle and vasculature	CVD	N/A
n-3 PUFA for Vascular Cognitive Aging	NCT01953705	A 3-year interventional study to investigate if omega 3 PUFA can support small blood vessels in the brain and promote brain health in adults >75 years of age with a high risk of cognitive decline	CVD	2
Impact of Ageing on Adipose, Muscle and Systemic Inflammation	NCT02777138	Observational study of macrophage and T-cells in adipose tissue and inflammatory markers (mRNA and protein secretions) between a group of younger and older males with similar lifestyles	CVD, T2D	N/A
Cell Signaling and Resistance to Oxidative Stress: Effects of Aging and Exercise	NCT03419988	An open interventional study investigating the effects of exercise on the expression of an antioxidant regulation protein nuclear erythroid-2-p45-related factor-2 (Nrf2) in peripheral blood mononuclear cells in healthy younger (18–28 years) and older (>60 years) adults	CVD, T2D	N/A
Resistance Exercise and Low-Intensity Physical Activity Breaks in Sedentary Time to Improve Muscle and Cardiometabolic Health (REALPA)	NCT03771417	An interventional study to assess the impact of resistance exercise with low-intensity physical activity breaks on skeletal muscle and cardiometabolic health in adults aged 65–80 years	CVD, T2D	N/A
Dietary Reduction of AGEs to Prevent Cognitive Decline in Elderly Diabetics	NCT02739971	An intervention pilot study to investigate if it is feasible to reduce dietary AGEs and therefore reduce cognitive decline	T2D	N/A
Inflammation				
Efficacy of Periodontal Treatment on Systemic Inflammation and for Prevention of Exacerbations in Patients With COPD (Expertention)	NCT03279718	An interventional pilot study to investigate if periodontal treatment can reduce systemic inflammatory markers (including CRP and IL-1b and IL-6)	COPD	N/A

Continued

TABLE 1 Continued

Study name	Study identifier	Study overview	Disease area	Phase
INvestigating COPD Outcomes, Genomics and Neutrophilic Inflammation With Tiotropium and Olodaterol (INCOGNITO)	NCT03152149	An open-label interventional trial comparing treatment of patients with COPD with tiotropium and olodaterol reduces bacterial load and neutrophilic inflammation <i>versus</i> inhaled fluticasone furoate and vilanterol	COPD	4
Pivotal Study to Assess the Efficacy, Safety and Tolerability of Dupilumab in Patients With Moderate-to-severe COPD With Type 2 Inflammation (BOREAS)	NCT03930732	An interventional study to assess the efficacy of dupilumab (an IL-4 receptor alpha antibody) in reducing the annual exacerbation rate in patients with moderate-to-severe COPD	COPD	3
Biological Effects of Quercetin in COPD	NCT03989271	A study to determine if a dietary supplement, quercetin, can enhance anti-inflammatory effects in patients with COPD and reduce markers of oxidative stress and inflammation	COPD	1/2
Anti-ST2 (MSTT1041A) in COPD (COPD-ST2OP)	NCT03615040	An interventional study to assess the use of an IL-33 inhibitor on the frequency of AECOPD events in patients with COPD over 48 weeks	COPD	2
A 12-week Study Treating Participants Who Have alpha1-antitrypsin-related COPD With Alvelestat (MPH966) or Placebo (ASTRAEUS)	NCT03636347	An interventional study investigating the effects of a neutrophil elastase inhibitor in patients with α_1 -antitrypsin deficient Pizz or null phenotype with COPD. Outcomes include sputum and blood inflammatory biomarkers, including neutrophil elastase activity.	COPD	2
Effect of IL-1 β Inhibition on Inflammation and Cardiovascular Risk	NCT02272946	An interventional study to investigate the effects of IL-1 β inhibitor canakinumab in reducing vascular inflammation in HIV-infected individuals	CVD	2
ASSessing the Effect of Anti-IL-6 Treatment in Myocardial Infarction: The ASSAIL-MI Trial (ASSAIL-MI)	NCT03004703	An interventional study to assess the impact of a single administration of an anti-IL-6 antibody, tocilizumab, on myocardial damage following myocardial infarction	CVD	2
Effects of SGLT-2 Inhibition on Myocardial Fibrosis and Inflammation as Assessed by Cardiac MRI in Patients With DM2	NCT03782259	An interventional study to investigate how inhibition of a glucose transporter, SGLT-2, with dapagliflozin impacts cardiovascular health and inflammation in patients with T2D	CVD, T2D	4
Cellular processes				
Early iNO for Oxidative Stress, Vascular Tone and Inflammation in Babies With Hypoxic Respiratory Failure	NCT01891500	An interventional study to investigate an already approved intervention, inhaled nitric oxide, in newborns with hypoxic respiratory failure to reduce biomarkers of oxidative injury	CVD	4
Impacts of Mitochondrial-targeted Antioxidant on Peripheral Artery Disease Patients	NCT03506633	An interventional study to examine the impact of a mitochondrial antioxidant (MitoQ) on vascular endothelial function in patients with peripheral vascular disease	CVD	N/A
Effects of Saxagliptin on Adipose Tissue Inflammation in Humans	NCT02285985	An interventional study to investigate the effects of saxagliptin on adipose tissue inflammation in obese participants	T2D	4

AECOPD: acute exacerbations of COPD; PUFA: polyunsaturated fatty acid; AGEs: advanced glycation end-products; CRP: C-reactive protein; IL: interleukin; MRI: magnetic resonance imaging; DM2: type-2 diabetes; N/A: not available; iNO: inhaled nitric oxide.

recognition that our diagnostic criteria may not identify early disease. For example, in COPD, approximately 70% of small airways are lost prior to the development of airflow obstruction; T2D is preceded with a period of “prediabetes” where blood sugars are raised, and atherosclerosis is usually only diagnosed when there is flow-limiting disease associated with symptoms. Identifying the predominant pathways in individual patients, while having the potential to provide therapeutic benefits, will require large, longitudinal and deeply phenotyped patient cohorts across disease and science silos.

Current studies of these mechanisms have often been conducted with one condition central to the study without considering the other diseases. Ongoing studies in each of the topics discussed in this article are summarised in table 1 and highlight the segregation of these studies, but also the lack of novel drugs and

the current focus on diet and exercise intervention. To understand these processes in more depth will require exacting science in carefully characterised patients with a cross-disciplinary approach.

It is also important to understand the relative contributions of each mechanism (ageing and senescence, inflammatory cytokines, altered cell processes) to each disease (COPD, CVD and T2D). The limited success of single cytokine therapies [129, 158–160], clearly demonstrates these diseases are complex and that each mechanism is just a piece of the puzzle. Identifying the relative contribution is challenging, as existing models usually allow either enough resolution to determine a single mechanistic effect, or identify broad changes without ascertaining the mechanism or the role of each disease in multimorbid patients. Addressing this challenge relies on a careful balance of *in vivo* models and *in vitro* human studies to stitch together the bigger picture, incorporating novel methods of network analysis and systems biology [161], and stratification of multimorbid patients in clinical studies, not just reporting them. This is further confounded by many studies not detailing the comorbidity of their patient populations; highlighted by one study that demonstrated the burden of comorbidity in COPD as of 8656 patients diagnosed with COPD, 1631 participants had an existing diagnosis of CVD or T2D and found associations with systemic inflammation and hospitalisation due to these comorbidities [162]. We might be missing important links in the mechanism of each disease because the multimorbid status of patients is not known or reported.

Conclusion

Finding treatments for multimorbidity is challenging but it is important to consider diseases holistically and not in isolation, as many pathological processes appear to be shared. There are multiple ways that COPD, atherosclerosis and T2D are linked, not just through shared risk factors but also through inflammation, ROS production, senescence and accelerated ageing and altered cellular functions and phenotypes. These processes are inter-related, each influencing the other and understanding the fine balance between positive cellular responses and damaging ones in health and across diseases may revolutionise therapy for chronic inflammatory diseases. Indeed, advancements in our understanding of chronic diseases are already suggesting new potential therapeutic strategies that bridge traditional disease silos; however, these are still in their infancy. The appreciation of neutrophil heterogeneity and the roles these cells play across chronic illness may also highlight potential treatment targets. Manipulation of T-cells and dendritic cells has yielded novel cancer therapies and the same concept might support new approaches for treating chronic inflammatory conditions [163, 164].

Most translational studies of COPD have focused on the lung disease and not fully characterised the burden of multimorbidity, but perhaps now is the time for a more thorough clinical assessment of the patients included in discovery science studies, detailing multimorbidity where it is present. It may be possible that systemic inflammation is simultaneously influencing the pathogenesis of COPD, atherosclerosis and T2D, yet limitations in diagnosis result in one being diagnosed first. Multimorbidity is also an important consideration for future clinical trials, as an intended target in one condition might improve an outcome for a related chronic disease, if we only had the foresight to look.

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