GRUPPO LINFOMI IN PAZIENTI CON IMMUNODEFICIT

Milano, UNAHOTELS Galles

23 maggio 2025

Come valutare l'inflammaging nella patogenesi dei linfomi e nella malattia di Castleman

Lara Gibellini

Nothing to disclose

Outline

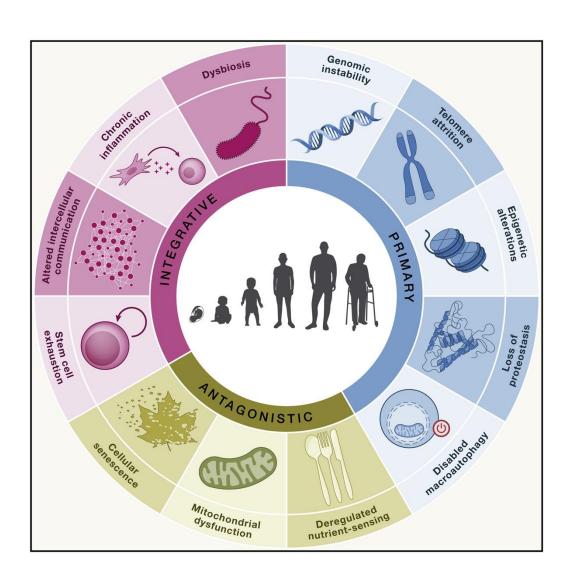
- Definition of inflammaging
- Possible causes
- Possible consequences
- How to measure inflammaging
- Considerations on lymphoma and Castleman Disease

Persistent chronic inflammation: inflammaging

- Low-grade, chronic state of inflammation
- Increased levels of IL-6, CRP, fibrinogen, IL-1β, TNF among others

• Still unknown whether inflammaging, which represents a risk factor for most age-related pathologies, is the cause or rather the effect of the aging process.

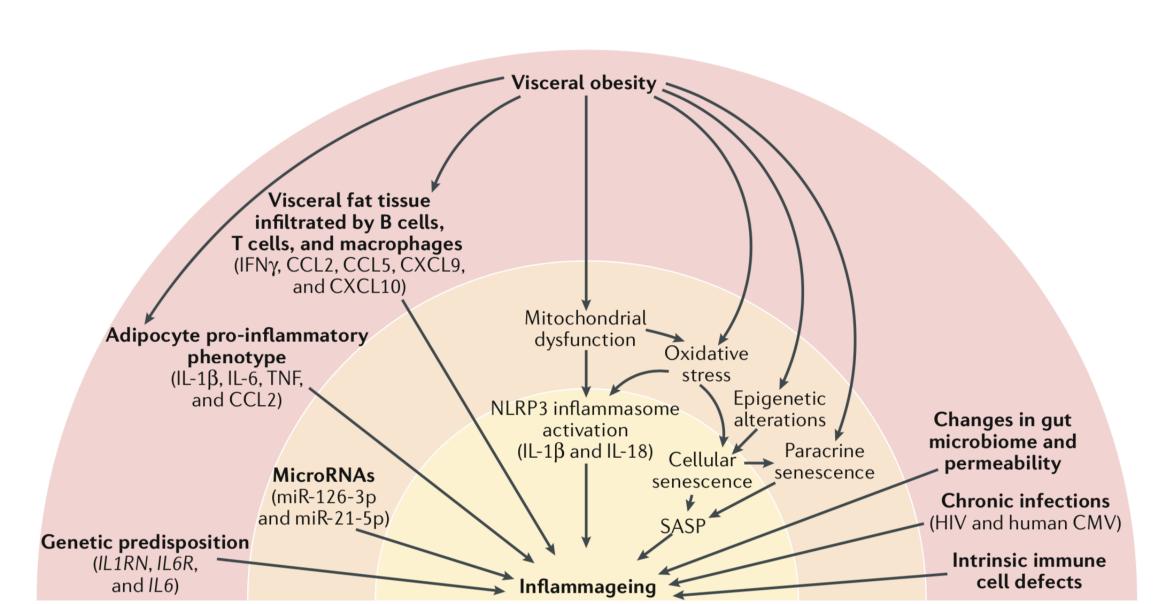
The hallmarks of aging

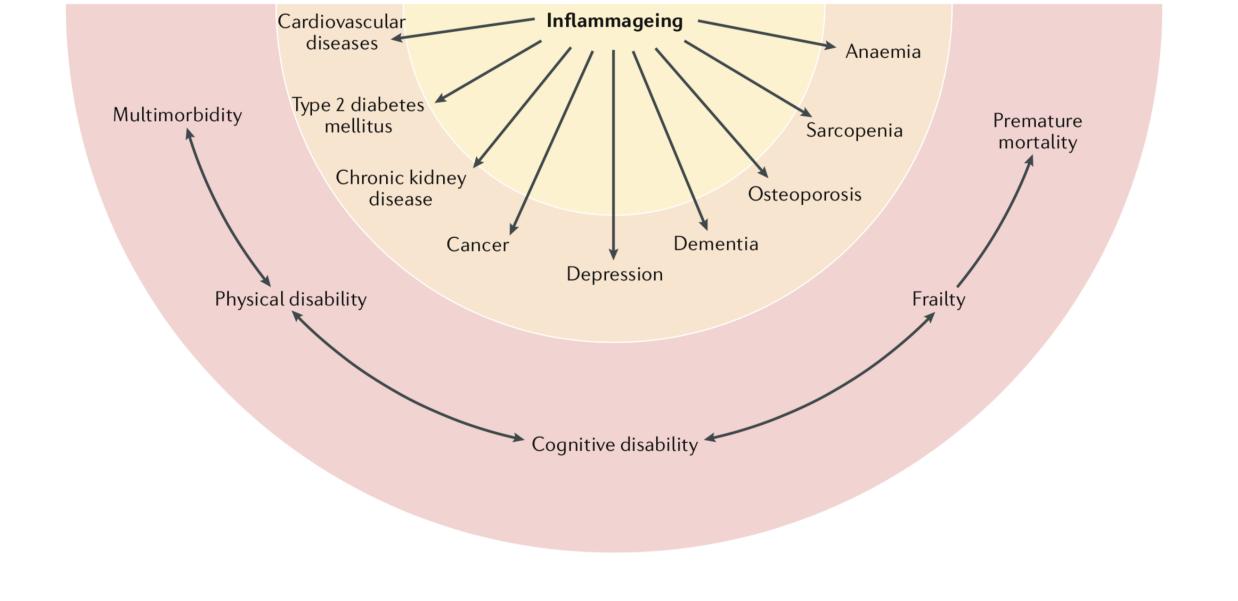


- Genomic instability
- Telomere attrition
- Epigenetic alterations
- Loss of proteostasis
- Disabled macroautophagy
- Deregulated nutrient-sensing
- Mitochondrial dysfunction
- Cellular senescence
- Stem cell exhaustion
- Altered intercellular communication
- Chronic inflammation
- Dysbiosis

Possible causes of inflammaging

Adapted from Ferrucci et al, Nat Rev, 2018



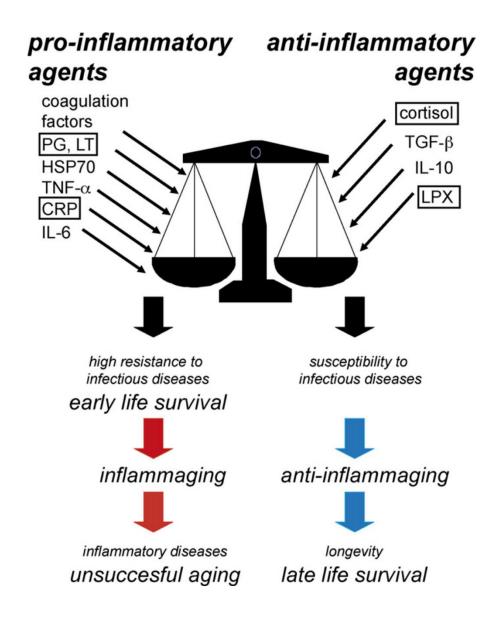


Possible consequences of inflammaging

Biomarker	Molecules involved	Effects
Pro-inflammatory and anti-inflammatory cytokines	TNF, IL-1 β , IL-6, IL-8, TGF- β , IL-10, CXCL9	Circulating biomarkers of chronic inflammation (IL-6 central role in inflammaging)
Other plasma soluble factors	CRP Fibrinogen	
N-linked glycan profile	Serum IgG-G0 digalactosylated or agalactosylated N-linked glycan structures	Biological age (pathological vs non-pathological ageing)
DNA methylation	353 CpG sites in ELOVL2 that wwere used to construct the epigenetic clock to estimate the methylation age	Chronological age and biological age
Circulating miRNA	miR-155 mIR-21 mIR-146a	Systemic inflammation
Metabolomics and lipidomics	Glycerophosphoethanolamines Glycerophosphocholines Glycerolipids Bile acids Steroids Isoprenoids Fatty amides Sphinolipids Trp levels L-carnitine esters	Healthy aging (centenarians)
Circulating cf-mtDNA	Cf-mtDNA	Systemic inflammation

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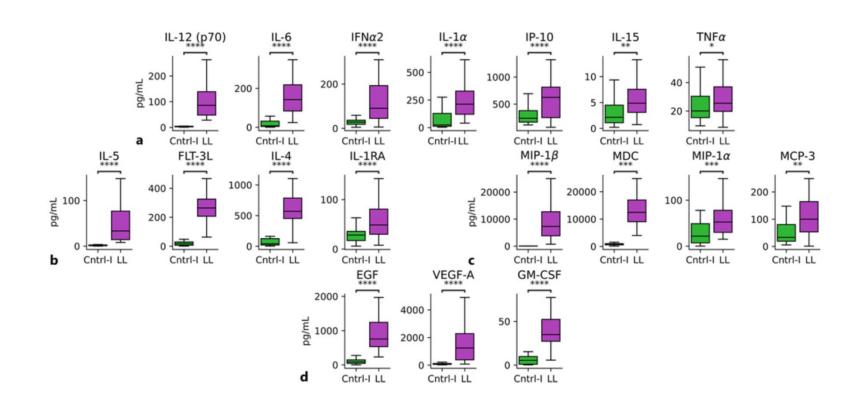
Inflammaging vs anti-inflammaging



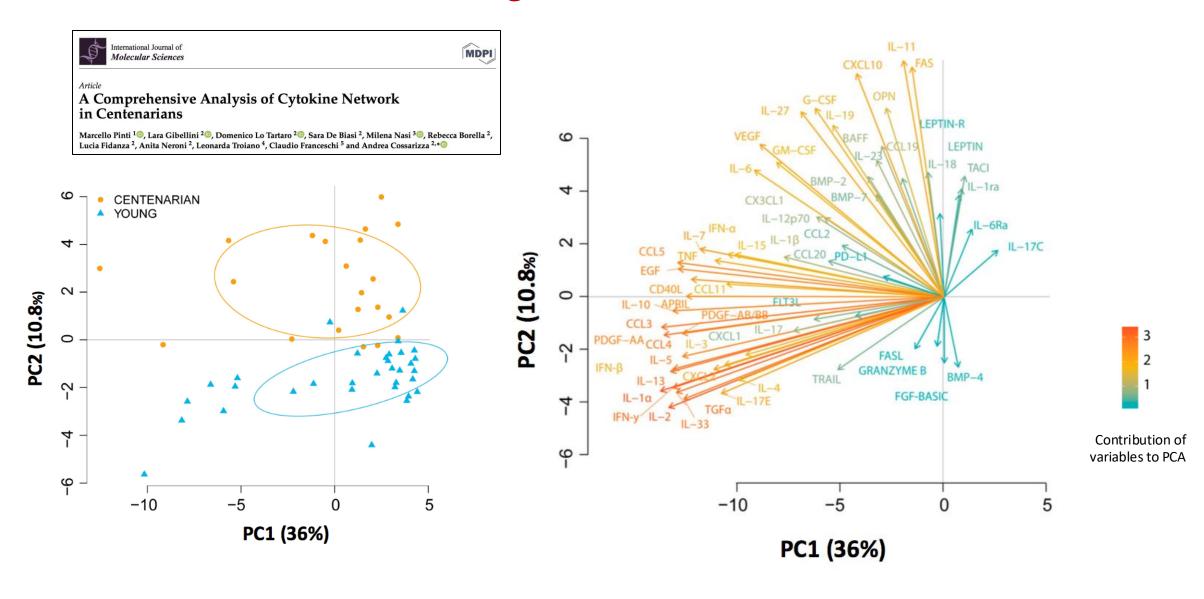
Inflammaging vs anti-inflammaging

The most longeve humans seem to be equipped with gene variants that allow them to optimize the balance between pro- and anti-inflammatory molecules.

Thus, their immune system is able to **minimize** the effects of the lifelong exposure to environmental insults and stressors.

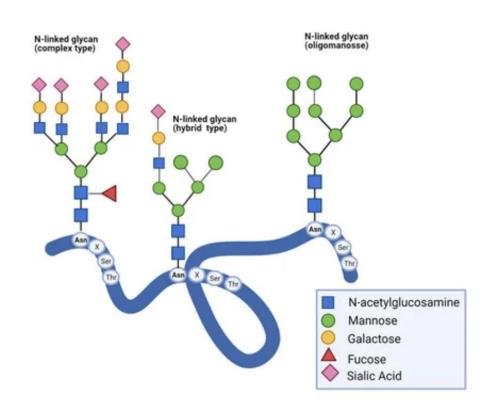


PCA of plasma cytokine dataset identifies groups of donors, *i.e.* middle-aged donors vs. centenarians



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N-glycans

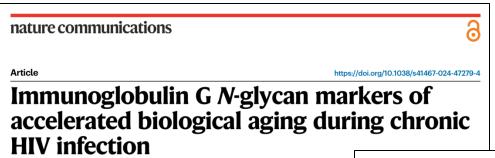


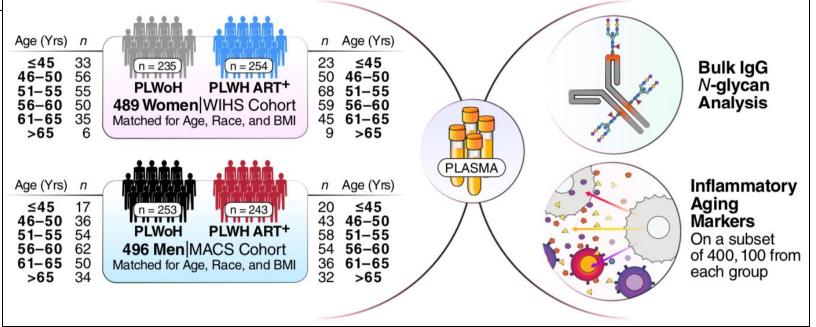
The log ratio of the relative abundance of two *N*-linked glycan species (namely, agalacto core-α-1,6-fucosylated diantennary glycan (NGA2F) and digalacto core-α-1,6-fucosylated diantennary glycan (NA2F)) increases progressively with age and is associated with features of healthy and unhealthy ageing.

N-glycans and inflammaging

- Protein galactosylation is responsible for the anti-inflammatory function of immunoglobulin G (IgG)
- With increasing age, the galactosylated biantennary structures that decorate the Asn297 of the crystallizable fragment (Fc) portion of IgG become devoid of galactose at both branches (called **IgG-G0**) and become highly pro-inflammatory.
- The study proposed to use IgG-G0 as a biomarker of inflammatory conditions during ageing, in which chronic low-grade inflammatory pathways negatively affect the glycosylation machinery of antibody-producing cells.

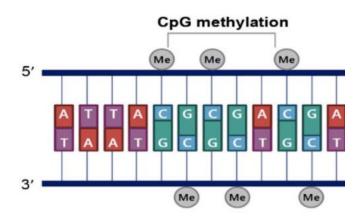
N-glycans markers of accelerated biological aging during chronic HIV infection



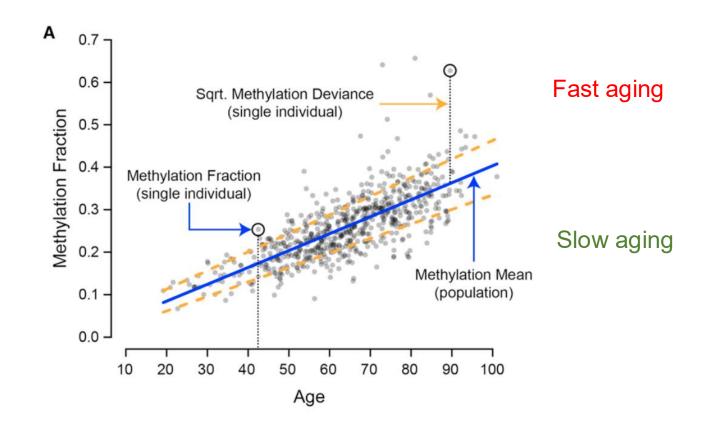


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Pro-inflammatory and anti-inflammatory cytokines	TNF, IL-1β, IL-6, IL-8, TGF-β, IL-10, CXCL9			
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Methylation profiles and human aging

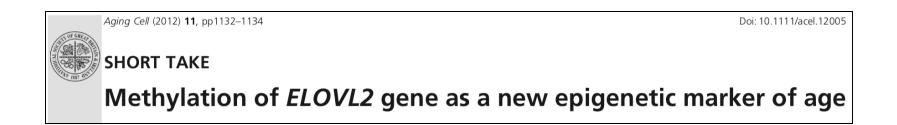


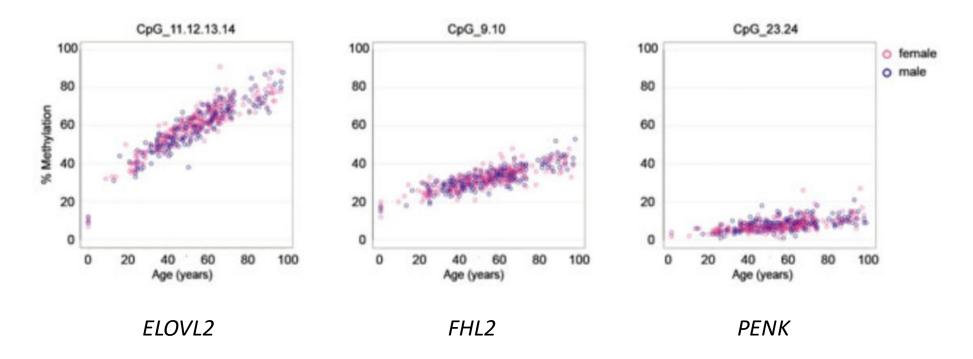
CpG methylation is a form of DNA modification where a methyl group is added to a cytosine base within a CpG dinucleotide



DNA mehylation is a biomarker of chronological aging and biological aging

Methylation profile of *ELOVL2* promoter





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Lipid and metabolic profiles and aging

- Centenarians had a peculiar lipid profile, with unique changes in 41 of 161 measured lipid species.
- The lipid profile emphasized that long-living individuals have marked features of anti-inflammatory molecules, such as increased levels of **phenylalanine**, which inhibits the nuclear factor-κB (NF-κB) pathway, and decreased levels of **glycerophosphocholine** (a circulating marker of cellular senescence). Monotliu et al, Aging, 2014
- Female familial longevity were identified. A profile that included high levels of phosphocholine and sphingomyelin and low levels of phosphoethanolamine and long-chain triglyceride species was found to be characteristic of healthy ageing. Gonzalez-Covarrubias et al, Aging Cell, 2013
- The plasma and urine from centenarians showed changes in the levels of specific glycerophospholipids and sphingolipids and a decrease in tryptophan concentration.
 Collino et al, Plos One, 2013.

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Mitochondria and inflammation

CLINICAL IMMUNOLOGY

Culprits with evolutionary ties

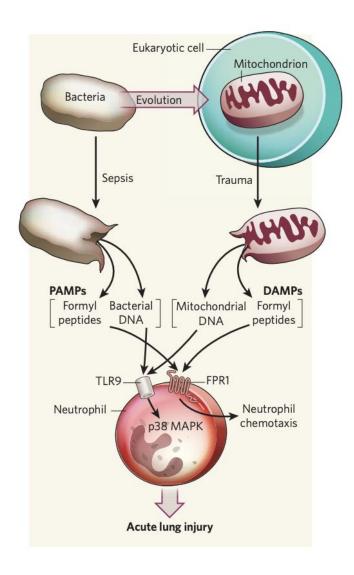
Carolyn S. Calfee and Michael A. Matthay

The cellular organelles we know as mitochondria are thought to have originated as symbiotic bacteria. Indeed, the two use common mechanisms to trigger innate immune responses to injury and infection, respectively.

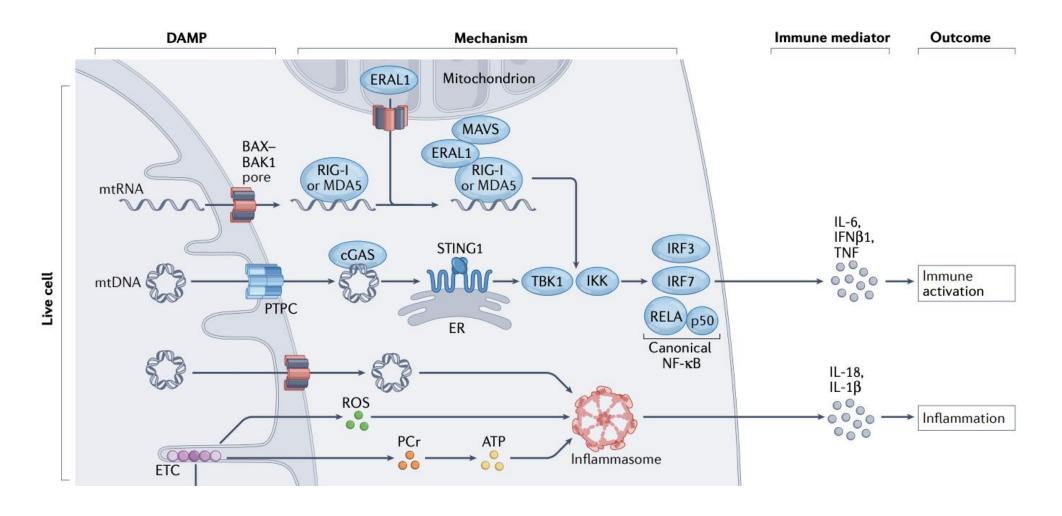
Mitochondria have originated as symbiotic bacteria



They use common mechanisms to trigger INNATE IMMUNE RESPONSES to injury and infection



Mitochondria and inflammation



mtDNA: DANGER SIGNAL

Plasma mtDNA increases with aging

Marcello Pinti et al.

DOI: 10.1002/eji.201343921

Eur. J. Immunol. 2014. 44: 1552-1562

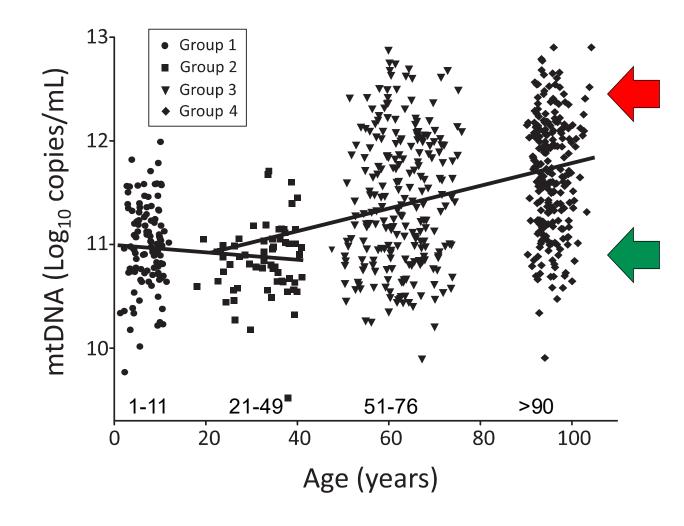


Circulating mitochondrial DNA increases with age and is a familiar trait: Implications for "inflamm-aging"

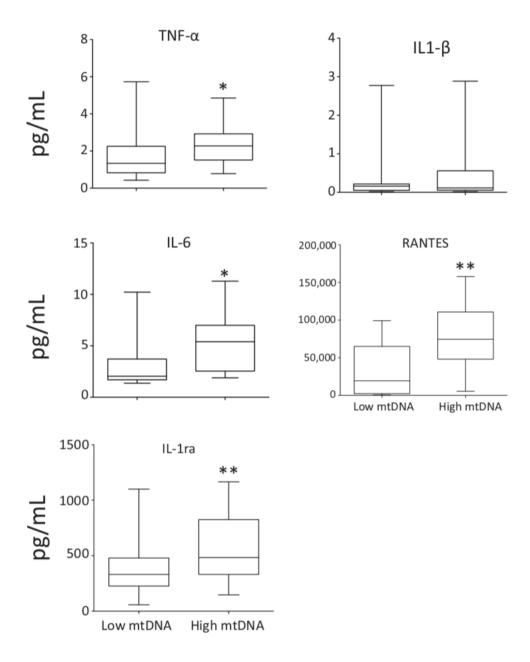
Marcello Pinti*¹, Elisa Cevenini*^{2,3}, Milena Nasi⁴, Sara De Biasi⁴, Stefano Salvioli^{2,3}, Daniela Monti⁵, Stefania Benatti¹, Lara Gibellini⁴, Rodolfo Cotichini^{6,7}, Maria Antonietta Stazi⁶, Tommaso Trenti⁸, Claudio Franceschi^{2,3} and Andrea Cossarizza⁴

Linear Regression for log 10 mt DNA by age

Groups	Number of obs.	R-squared	Beta Co eff.	Age p	959	% CI
Group 1 and 2	171	0.0215	-0.0045	0.055	-0.0091	0.0001
Group 2, 3 and 4	4 516	0.1590	0.0115	<0.001	0.0092	0.0138



Plasma mtDNA increases with aging



Lymphoma and inflammaging

Chronic immune activation

Persistent inflammation can cause DNA damage and promote mutations in lymphocytes.

Immunosenescence

Aging impairs immune surveillance, making it easier for malignant clones to escape detection.

Microenvironmental changes

Inflammaging alters the tissue microenvironment, which may support lymphoma growth and survival.

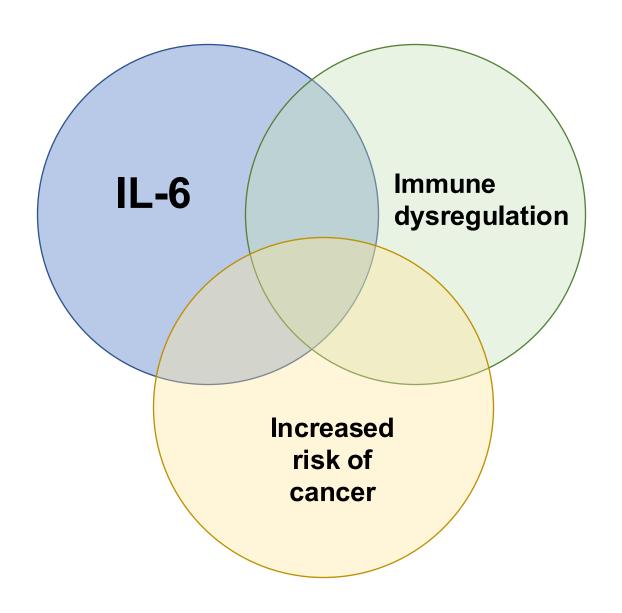
Cytokine imbalance

Pro-inflammatory cytokines involved in inflammaging may contribute to the growth and survival of lymphoma cells.

Clinical Implications

Age-adjusted treatment strategies are critical in older lymphoma patients, who may also be affected by inflammaging-related frailty.

Castelman disease and inflammaging



Conclusions

- Low-grade, chronic state of inflammation
- Inflammaging has multiple and heterogeneous causes
- Inflammaging has multiple and heterogeneous consequences
- Inflammaging can be measured
- There are several links between inflammaging and lymphoma, and between inflammaging and Castelman disease

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