Gender and longevity
Rita Ostan1, Daniela Monti2, Claudio Franceschi1

Summary. At present, women show a higher life expectancy than men and this gender difference is becoming a world-wide phenomenon indicating the central demographic role of gender in ageing and longevity. A combination of genetic, environmental, historical, anthropological, socio-economic and cultural aspects as well as the geographical origin is at the basis of the longer survival of women. Maternal transmission is determining in all the three pillars of human genetics of ageing and longevity (nuclear genetics, mitochondria genetics, and microbiome genetics). Moreover, mothers and grandmothers play a key role in the offspring and grandchildren care/survival by contributing to their wellbeing and transmitting traditional and hygienic habits able to modulate life-long health status and lifespan. Finally, women live longer than men but they show a worse quality of life in advanced age indicating that gender is probably the most significant variable for the lifelong health status.

Introduction
Among humans, it is well known that at present women live longer than men, and gender difference in life expectancy is becoming a world-wide phenomenon both in industrialized nations and in low-income countries. Therefore, gender plays a central demographic role in ageing and longevity. In the biology of ageing the impact of gender difference has been extensively evaluated, but it still remains largely unexplored territory, particularly regarding the interconnection between a series of fundamental aspects such as hormonal, immunological and metabolic pathways. All these themes will be extensively addressed in the next issues of this journal. In the current article, we will concentrate on topics that have generally received less attention in previous literature, and we will focus on some genetic and cultural specificities regarding women in order to help explain some of the reasons that allow them to live longer.

It is now clear that life span as well as longevity are complex and multifactorial traits resulting from an intriguing combination of “Nature” and “Nurture”, the unique reciprocal interaction between genetics and environment which characterizes every human being. Thus, we can start from the assumption that a peculiar combination of genetic, environmental, historical, anthropological, socio-economic and cultural factors as well as geographical origin contributes to longer female life expectancy worldwide.

Genetic aspects of longevity

The complex role of the double X chromosome in ageing and longevity

In the mammalian cells of a young woman, one of the two X chromosomes is epigenetically and randomly inactivated in early embryonic life. Females are therefore a mosaic of cells in which either the maternal or the paternal X chromosome is inactivated1, and the ratio is close to 50% for each chromosome. Several studies have shown that skewed X chromosome inactivation (XCI) occurs with age in blood cells2,3. We investigated this phenomenon, employing an experimental model of longevity/healthy ageing consisting

1. Department of Experimental, Diagnostic and Specialty Medicine (DIMES) and Interdepartmental Centre “L Galvani” (CIG), University of Bologna, Italy; 2. Department of Clinical, Experimental and Biomedical Sciences, University of Florence, Florence, Italy.

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of female centenarians, their female offspring, sex- and age-matched controls (offspring born from shorter-lived parents) and young women. The model of centenarians and their offspring has allowed us to demonstrate the better health status and significant survival advantage of centenarian offspring, i.e. reduced prevalence of several age-associated diseases (cardiovascular, respiratory, osteoarticular diseases and cancer) and improved functional fitness (handgrip strength, chair stand, etc.)4. We confirmed that XCI skewing is an age-dependent phenomenon and demonstrated that this process was significantly less severe and frequent in centenarian offspring compared to their age-matched controls5. The underpinning mechanisms and biological consequences of XCI skewing with age still remain unclear. Within this scenario, our results highlight a possible detrimental link between the rate of XCI skewing and healthy ageing/longevity, fitting the hypothesis that the female mosaic is a winning strategy, sustaining a cooperative adaptive mechanism with possible advantages in pathological and physiological conditions. Conversely, the absence of a similar mosaic strategy in men might contribute to their shorter lifespan6,7.

Mitochondria, ageing and longevity: the case of fibroblasts from centenarians

Mitochondria are fundamental organelles for cellular energy/ATP production and are actively involved in a variety of cellular processes, including urea cycle, heat production, apoptosis, inflammasome activation and cell senescence, among others. Mitochondria are also the main producers of reactive oxygen species (ROS), the most important by-product of OXPHOS. It has been suggested that, besides their physiological role in cell signalling, they may play a role in the ageing process as well as in age-related diseases. Data from primary culture of fibroblasts from long-lived individuals, including female centenarians, indicated that such cells displayed a lower complex I-driven ATP synthesis and a higher production of H₂O₂ compared with fibroblasts from old and young subjects. However, the bioenergetics of centenarian cells was well preserved and similar to that of younger subjects owing to a compensatory response provided by increased mitochondrial mass and hyperperfused and elongated mitochondria. Elongated mitochondria, such as those found in centenarian fibroblasts, are largely spared from autophagic degradation, show increased levels of dimerization and ATP synthase activity, and maintain ATP production more efficiently than fragmented mitochondria. This longevity-associated mitochondria “hypertrophy” can be considered a “remodelling” mechanism capable of compensating for age-related functional defects and preserving bioenergetic functionality even at an extreme age, which likely contributes to longevity8.

The intriguing role of maternally-inherited mitochondrial DNA (mtDNA) in human longevity

In the context of complex biological role of mitochondria, mitochondrial DNA (mtDNA) variability represents another aspect which merits particular attention in the study of longevity. Human mtDNA is a constitutive part of the genetic machinery of each cell, within the framework of continuous crosstalk between mtDNA and the nuclear genome. mtDNA (16,569 bp) encodes a few qualitatively and quantitatively relevant genes, being a constitutive component of OXPHOS and owing to the high copy number of mtDNA in each cell. mtDNA is only inherited through the mother and its germline variants (haplogroups) have been found to be associated with longevity in several populations, suggesting the presence of a maternal component of human longevity. In particular, the large EU project GEHA (Genetics of Healthy Ageing) studied 2,200 ultra-nonagenarian (90+) from different European countries belonging to 90+ sib-pairs together with the same number of sex- and geographically-matched younger controls, and was able to identify different haplogroups related to longevity in both males and females. The J2 haplogroup was associated with male longevity while H2 and T2 haplogroups were associated with female longevity9. Taking advantage of the complete sequencing of a high number of mtDNA molecules, it was also possible to evaluate for the first time the cumulative effect of specific and concomitant mtDNA mutations, including those that have a low or very low impact per se. The analysis of the mutations occurring in different OXPHOS complexes showed an intricate scenario with a different mutation burden in nonagenarian subjects versus controls. In particular, mutations in subunits of OXPHOS complex I had a beneficial effect on longevity, while the simultaneous presence of mutations in complexes I and III and in complexes I and V appeared to be detrimental9. The final conclusion was that “particular rare mtDNA mutations present only in specific populations might be beneficial (or detrimental) for longevity and may explain part of the genetic component of longevity in that population, similarly to what has been suggested for private nuclear DNA polymorphisms”9.

MtDNA mutations are transmitted from centenarian mothers to their progeny

One of the factors that can contribute to ageing and longevity is the accumulation with age of mtDNA mutations. mtDNA heteroplasmy, i.e. the presence in
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potentially pathogenic bacteria (pathobionts) with respect to beneficial ones (symbionts producing butyrate and other short chain fatty acids)\(^1\).

On the whole, these considerations indicate that genetics of human ageing and longevity is dependent on three pillars (nuclear genetics, mitochondria genetics, and microbiome genetics)\(^2\) that must be evaluated together. It is worth noting that maternal transmission plays a key role in all three areas of genetics and in their relationship with human longevity.

**Maternal cultural habits, offspring survival and longevity**

Socio-economic and cultural/anthropological components are fundamental contributors to human longevity and are likely related to offspring care/protection as well as transmission of behavioural habits whose influence can have a lifelong duration. A woman who protects her own health and survival through her behaviour will also protect the health of her offspring, both directly (as she is in a better state of health at the time of childbearing) and indirectly (as she is more mindful about her children’s health)\(^3\).

Healthy ageing could be advantageous and therefore positively selected for, if parents can influence improved fertility and/or survival of their children, grandchildren and/or other family members. The “Grandmother hypothesis” argues that caregiving by older women who underwent menopause could improve grandchildren’s survival by alleviating the child burden of their daughters, who in turn could have more successful/numerous pregnancies and offspring. Therefore, mothers and grandmothers have assumed and continue to play a key role in the care/survival of offspring and grandchildren, by contributing to their health, by providing nutrition and by passing on traditional habits and hygienic guidelines. These practices and life-style behaviours have lifelong effects starting from intrauterine life and may modulate health status and lifespan in advanced age. However, it is also important to consider that mothers and grandmothers may also have a negative influence on offspring (by not breastfeeding, smoking and drinking alcohol, drug abuse, unbalanced diets, among other factors).

In any case, mothers do have the role and responsibility of providing the first nine months of environment to human beings. A uterine environment is sensitive to any type of metabolic, infectious and stressful event affecting the mothers, and is now recognized to be developmentally crucial in establishing a variety of set points (metabolic and epigenetic among others) in the offspring, which in turn may have long-term biological effects, as predicted by the hypothesis of the foetal origin of many diseases in adulthood and old age.

**Conclusions**

All the existing data suggests that one of the most powerful strategies for reaching exceptional longevity is to be female. Women live longer than men because they die at lower rates\(^4\), but they pay for their survival advantage with a worse quality of life due to the increased prevalence of a variety of non-lethal pathological conditions. Old women display a higher overall rate of physical disability and are more likely than men to report difficulties in walking, climbing stairs and other common activities\(^5\). This gender dichotomy in ageing is confirmed by the centenarian model, of which females form the majority, but are characterized by poorer health as well as more impaired functional and cognitive status indicating that, even in the last phase of human life, gender is probably the most important variable of lifelong health status. This final consideration is in our opinion the most robust scientific justification for starting a new journal specifically focused on gender-specific medicine, an emerging, current and somewhat neglected topic which we are beginning to appreciate in its fascinating complexity and which deserves much more attention than we previously thought.

**References**


