

EDITORIAL COMMENT

# Telomeres and Atherosclerosis

## The Attrition of an Attractive Hypothesis\*

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In this issue of the *Journal*, the paper by Fernández-Alvira et al. (1) is probably the (pen)ultimate nail in the coffin of the attractive hypothesis that cross-sectionally measured leukocyte telomere length (LTL) is causally involved in the development of atherosclerosis. In this study of 1,459 middle-aged cardiovascular disease (CVD)-free men and women, neither average LTL nor short telomere load was a significant independent predictor for the presence of subclinical atherosclerosis. These findings are in line with previously published reports from the Asklepios and Bruneck cohorts, which similarly did not find an association between average LTL and early atherosclerosis (2,3).

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Importantly, the current study extends these prior findings in 3 important ways. First, the phenotype of atherosclerotic burden is more extensive and, in all likelihood, also more precise than in previous studies. Subclinical atherosclerosis was evaluated by collating data from a coronary artery calcium scan, combined with 2-dimensional ultrasound of the infrarenal aorta and iliofemoral arteries and 3-dimensional ultrasound of the carotid and femoral arteries. Moving from binary “plaque presence” toward a more granular “plaque burden” is an important step forward. Second, LTL was measured with quantitative fluorescence in situ hybridization, instead of the previously used Southern

blot (2) or quantitative polymerase chain reaction (3) techniques. Although quantitative fluorescence in situ hybridization is not necessarily more accurate, it was at least sufficiently sensitive to detect known associations among LTL and age, sex, and oxidized low-density lipoprotein serum levels in this cohort. Current results, therefore, pinpoint biology rather than methodology to explain the lack of a significant association between LTL and early atherosclerosis. Third, previous studies looked at average LTL and atherosclerosis. The findings from the PESA (Progression of Early Subclinical Atherosclerosis) study extended this by an insightful subanalysis investigating the association between atherosclerosis and the load of short LTL (<3 kb), which also proved negative.

To understand the wider relevance of this paper, it is not enough to merely look at what this paper adds to the field of telomere biology. The ultimate importance is closely related to understanding what makes the study of telomere biology potentially so attractive to a cardiologist.

The true driving force behind interest in telomere biology is the tantalizing promise of a biomarker that captures aging by integrating lifelong exposure to risk factors, including those that are known, unknown, and misunderstood, and thereby reflects the cumulative burden of subclinical damage. Such a biomarker should be far more informative in providing risk stratification and potential for therapeutic guidance than a biomarker that merely reflects a single point in time (4).

Both imaging-derived (intima-media thickness, plaques, and coronary calcium scores) and, even more so, biomechanics-derived time-integrative biomarkers (arterial stiffness, reflection magnitude, and pulse wave velocity) have, to a varying degree, proven additive value (5-8). Still, a time-integrative biomarker, obtainable from a simple blood draw (such as LTL), would be more easily applicable for prevention at the population level.

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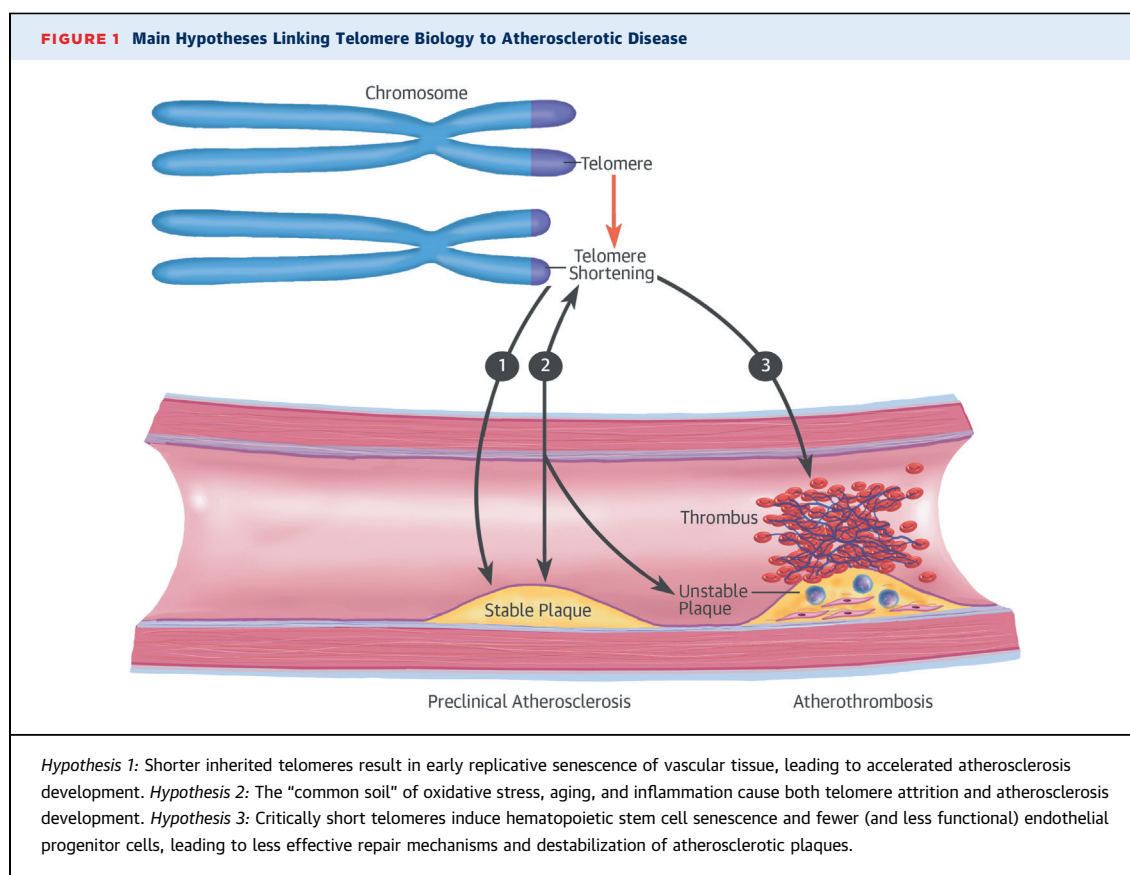
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Telomeres are TTAGGG-hexamer repeat containing nucleoprotein complexes that protect the chromosomal termini by formation of a complex loop structure. A subject's telomere length is largely inherited (although only in part genetically) and is very similar between tissues. In somatic tissues, telomeres are characterized by age-dependent attrition due to cell division and oxidative stress, which may ultimately lead to replicative senescence (one of Nature's primary defense mechanisms against cancer development) (9,10).

Widespread interest in a potential link between telomeres and CVD was accelerated by the publication by Samani et al. (11) in 2001 showing shorter average LTL in subjects with advanced coronary artery disease compared with disease-free control subjects, a finding that was confirmed in a recent meta-analysis (12). Furthermore, results from genome-wide association and prospective studies suggest causality with CVD (13,14). A recurrent hypothesis therefore stated that inherited shorter telomeres result in early replicative senescence of vascular tissue, thereby inducing atherosclerosis and predestining subjects to future CVD. Yet, if LTL shortening is causally related to CVD

through atherosclerosis initiation, it is highly paradoxical that findings from the PESA study and others show no association with early (pre-clinical) atherosclerosis (1,2).

Other potential causal and noncausal mechanisms linking telomeres to CVD have been postulated (Figure 1). One school of thought considers shorter telomeres as a pure epiphenomenon, the result of accelerated telomere shortening due to a cluster of shared risk factors also involved in atherogenesis, such as hypertension, oxidative stress, and inflammation (15-17). An extension of this hypothesis suggests that the resulting shorter telomeres may themselves have a causal effect in a much later stage of the atherosclerotic process (16), although it should be noted that the total burden of attrition on adult LTL is very limited compared with the overall effect of inheritance (18). As both atherosclerosis and LTL reflect cumulative exposure to partially shared risk factors over time, there is a high likelihood of a spurious association. This "associative bias" will be reinforced as longer timeframes and more advanced disease are considered and is likely to survive statistics adjusting for confounding risk



factors, which are usually measured at only 1 point in time.

A final alternative hypothesis assuming causality focuses on the transition from silent atherosclerosis to a cardiovascular event. For example, critically short telomeres could induce hematopoietic stem cell senescence and fewer (and/or less functional) circulating endothelial progenitor cells, which leads to less effective endothelial repair and subsequent destabilization of atherosclerotic lesions, triggering cardiovascular events (19). Interestingly, this hypothesis bridges the conflicting associations with late-stage diseases in the absence of an association with pre-clinical disease (16).

Taken together, current evidence suggests that although there is an association between LTL and atherothrombotic cardiovascular events, there is

no association with pre-clinical atherosclerosis. The findings from the PESA study confirm and extend previous findings by adding both a very performant phenotyping of pre-clinical atherosclerosis at the population level and an alternative, yet innovative, methodology to study the more physiologically relevant short telomere load. Still, the potential of a potent “early vascular aging” biomarker available through a simple blood test is so alluring that research on telomere biology in the field of CVD is unlikely to end here.

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## REFERENCES

1. Fernández-Alvira JM, Fuster V, Dorado B, et al. Short telomere load, telomere length, and sub-clinical atherosclerosis: the PESA study. *J Am Coll Cardiol* 2016;67:2467-76.
2. De Meyer T, Rietzschel ER, De Buyzere ML, et al., for the Asklepios Study Investigators. Systemic telomere length and preclinical atherosclerosis: the Asklepios Study. *Eur Heart J* 2009;30:3074-81.
3. Willeit P, Willeit J, Brandstätter A, et al. Cellular aging reflected by leukocyte telomere length predicts advanced atherosclerosis and cardiovascular disease risk. *Arterioscler Thromb Vasc Biol* 2010;30:1649-56.
4. Nilsson PM, Boutouyrie P, Cunha P, et al. Early vascular ageing in translation: from laboratory investigations to clinical applications in cardiovascular prevention. *J Hypertens* 2013;31:1517-26.
5. Ben-Shlomo Y, Spears M, Boustred C, et al. Aortic pulse wave velocity improves cardiovascular event prediction: an individual participant meta-analysis of prospective observational data from 17,635 subjects. *J Am Coll Cardiol* 2014;63:636-46.
6. Chirinos JA, Kips JG, Jacobs DR Jr., et al. Arterial wave reflections and incident cardiovascular events and heart failure: MESA (Multiethnic Study of Atherosclerosis). *J Am Coll Cardiol* 2012;60:2170-7.
7. van Sloten TT, Sedaghat S, Laurent S, et al. Carotid stiffness is associated with incident stroke: a systematic review and individual participant data meta-analysis. *J Am Coll Cardiol* 2015;66:2116-25.
8. Den Ruijter HM, Peters SAE, Anderson TJ, et al. Common carotid intima-media thickness measurements in cardiovascular risk prediction: a meta-analysis. *JAMA* 2012;308:796-803.
9. De Meyer T, Vandepitte K, Denil S, et al. A non-genetic, epigenetic-like mechanism of telomere length inheritance? *Eur J Hum Genet* 2014;22:10-1.
10. Aviv A, Kark JD, Susser E. Telomeres, atherosclerosis, and human longevity. *Epidemiology* 2015;26:295-9.
11. Samani NJ, Boulby R, Butler R, et al. Telomere shortening in atherosclerosis. *Lancet* 2001;358:472-3.
12. D'Mello MJ, Ross SA, Briel M, et al. Association between shortened leukocyte telomere length and cardiometabolic outcomes: systematic review and meta-analysis. *Circ Cardiovasc Genet* 2015;8:82-90.
13. Brouillette SW, Moore JS, McMahon AD, et al., for the West of Scotland Coronary Prevention Study Group. Telomere length, risk of coronary heart disease, and statin treatment in the West of Scotland Primary Prevention Study: a nested case-control study. *Lancet* 2007;369:107-14.
14. Codd V, Nelson CP, Albrecht E, et al. Identification of seven loci affecting mean telomere length and their association with disease. *Nat Genet* 2013;45:422-7, 427e1-2.
15. Bekaert S, De Meyer T, Rietzschel ER, et al. Telomere length and cardiovascular risk factors in a middle-aged population free of overt cardiovascular disease. *Aging Cell* 2007;6:639-47.
16. De Meyer T, Rietzschel ER, De Buyzere ML, et al. Telomere length and cardiovascular aging: the means to the ends? *Ageing Res Rev* 2011;10:297-303.
17. Benetos A, Okuda K, Lajemi M, et al. Telomere length as an indicator of biological aging: the gender effect and relation with pulse pressure and pulse wave velocity. *Hypertension* 2001;37:381-5.
18. Benetos A, Kark JD, Susser E, et al. Tracking and fixed ranking of leukocyte telomere length across the adult life course. *Aging Cell* 2013;12:615-21.
19. Minamino T, Miyauchi H, Yoshida T, et al. Endothelial cell senescence in human atherosclerosis: role of telomere in endothelial dysfunction. *Circulation* 2002;105:1541-4.

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