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Issue: *Innate Inflammation and Stroke***Pharmacological modulation of vascular inflammation in atherothrombosis**Raffaele De Caterina,^{1,2} Marika Massaro,³ Egeria Scoditti,³ and Maria Annunziata Carluccio³¹Institute of Cardiology, "G. d'Annunzio" University, Chieti, Italy. ²Fondazione "G. Monasterio," Pisa, Italy. ³CNR Institute of Clinical Physiology, Lecce, Italy

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Vascular inflammation, especially at the level of endothelial cells, has been shown to play a pivotal role in the inception, progression, and clinical complications of atherosclerosis. The common denominators for the activation of inflammatory genes appear to be a small subset of transcription factors—among which include nuclear factor- κ B, activator protein-1 (AP-1), and GATA—that function as the central hub of vascular inflammation. Strategies directed to inhibit both the secondary mediators and the primary triggers (atherosclerosis risk factors) appear viable to inhibit atherosclerosis. However, attempts have now been made to address the central hub of vascular inflammation. "Old" drugs, such as dipyridamole, can also now be revisited for properties related to inhibition of vascular inflammation, probably by acting on the common hub of inflammation.

Keywords: inflammation; endothelial cells; endothelial activation; nuclear factor- κ B; dipyridamole

Introduction

Our knowledge on the pathophysiology of myocardial infarction and of atherosclerotic vascular disease in general has gradually accrued over the past two centuries but has received an important acceleration in the past 20 years. While myocardial infarction, since its first descriptions, has been linked to coronary artery disease, the instabilization of an otherwise smoldering disease—leading to acute coronary syndromes—has been only the product of knowledge of recent years. Today, "inflammation," in its broader sense, can comprehensively account for the pathogenesis of vascular disease, including its late clinical manifestations. Here, we will briefly review recent knowledge in this area and highlight how even "old fashioned" drugs may need a modern revisitation for properties previously unsuspected, ultimately yielding potential new tools to combat the disease of the past and the present century.

On the origin of myocardial infarction and arterial instabilization

About 40 years ago, knowledge on the origin and pathophysiology of atherosclerosis mostly revolved

on the entry of cholesterol esters into vascular intima and on the correlation of such accumulation with hemodynamic stress. Several studies had already indeed correlated high levels of blood cholesterol with atherosclerotic vascular disease. Other studies had correlated the specific location of atherosclerotic plaques in bifurcation and branching points with the shear stress caused by the turbulence of blood flow, considered responsible for the increased endothelial permeability.¹⁻³ The transport of cholesterol in a free form through endothelial cells and its deposition in intimal phagocytes was being recognized, and gradually the role of low-density lipoprotein (LDL) cholesterol gradually emerged as the principal trigger of atherosclerosis.⁴⁻⁶ The body of knowledge acquired in those years offered a reasonable interpretation on the origin and progression of atherosclerosis in relation to the well-appreciated risk factors for ischemic heart disease. But those years were also witnessing a dispute, now resolved and forgotten by most, on the pathogenesis of myocardial infarction on an atherosclerotic substrate: important schools of thought, on both sides of the Atlantic Ocean, considered the progressive severe stenosis of

atherosclerotic arteries as the final trigger for acute events.^{7,8}

Many at that time believed that thrombosis was secondary to blood stasis following the infarction,⁹ therefore a consequence and not a cause. Much of this knowledge was based on pathological sections of coronary arteries in autopsy material, where—retrospectively interpreted—the variable time lag between the onset of symptoms and death could account for the entire spectrum of possible findings, supporting both the causality and the noncausality hypotheses. If one study has to be quoted as a divider between the time of quarreling and the time of data consolidation indicating thrombosis as the final trigger of plaque instabilization, this is the study by De Wood *et al.*, which showed the greater incidence of total or subtotal coronary occlusion as an inverse function of time elapsed between symptom onset and the coronary angiography used to demonstrate the occlusion, with the retrieval of thrombotic material through a Fogarty catheter when such intervention was performed early in time.¹⁰ Davies and coworkers, in 1985, illustrated the concept of plaque fissuring as a common cause of the acute manifestations of ischemic syndromes.¹¹ Closing the causality loop were, a few years later, the clinical trials demonstrating the possibility of dramatically reducing postmyocardial infarction mortality with antithrombotic treatments.^{12–25} The focus at that point could shift upstream, to understand the precipitating causes of arterial thrombosis.

The inflammatory theory of atherosclerosis

Atherosclerotic lesions begin as fatty streaks, areas of intimal thickening due to the accumulation of “foam cells” (essentially lipid-laden macrophages), extracellular matrix, and lymphocytes.^{26–30} Mature plaques, instead, consist of a thickened intima containing a lipid-rich core (*atheros*, gruel), surrounded by extracellular matrix. The lipid-rich core contains abundant tissue factor, capable of activating the extrinsic pathway of coagulation.^{31,32} The presence of a collagen-rich cap and of smooth muscle cells around the core prevent the contact of blood with the thrombogenic component.³³ Smooth muscle cells, stimulated by the presence of transforming growth factor- β (TGF- β), fibroblast growth factor-2, (FGF2), and platelet-derived growth factor (PDGF) produced by cells participating in the inflammatory atherosclerotic process, indeed mi-

grate from the media towards the intima, secreting several types of collagen and extracellular matrix (types I and II collagen, elastin, proteoglycans, fibronectin), contributing to the resistance to rupture of the fibrous cap and preventing thrombosis,^{34–36} and switching from purely contractile cells into secreting and proliferating cells.³⁷ Matrix digestion instead occurs because of the production of specific proteinases, named matrix metalloproteinases (MMPs), mostly by activated macrophages arrived in the intima as transformation of circulating monocytes and their local replication.

Many such cellular aspects of atherosclerosis, such as the passage of monocytes from peripheral blood to tissues and macrophage activation, can be generically defined as “inflammatory.”^{32,38–40} Inflammation indeed, in its widest acception, can be defined as the passage of cellular elements from the blood to tissues in response to an injury on the vascular wall or of surrounding tissues.⁴¹ Acute inflammation (with the prevailing involvement of neutrophils), cell-mediated immune responses (with a prevailing involvement of lymphocytes), and atherosclerosis (with the primary involvement of monocytes) are three forms of “inflammatory” reactions, all characterized by phenomena of passage of cells from the blood to the tissues. Inflammatory aspects are also the secretion of matrix and the collagenolytic degradation of the matrix by enzymes, mostly deriving from macrophages. Inflammation is therefore at the basis of all pathogenetic aspects of atherosclerosis, from lesion inception (the adhesion and migration of monocytes),^{41,42} to their growth (accumulation of cells and matrix),⁴³ up to their clinical emergence (plaque fissuring).⁴⁴

Many molecular aspects of vascular inflammation present in atherosclerosis are also common to acute inflammation and immune phenomena. All these have at their basis a condition of endothelial dysfunction (endothelial “activation”) with the expression by the endothelium of an adhesive phenotype towards various leukocyte subtypes from peripheral blood.⁴¹ The very same localization of atherosclerosis at particular sites of the vascular tree can be explained by such phenomena. Indeed, the preferential localization of atherosclerosis in branching sites, where shear stresses are low or oscillating, and produce temporal fluctuations in the tangential friction exerted on endothelial cells,^{45–47} can be explained by the activation

of specific receptors responsive to such forces (shear stress response sensors) and the ensuing nuclear events involving shear stress response elements (SSRE). These produce a cytoskeletal rearrangements of endothelial cells, the surface expression of endothelial adhesion molecules (intercellular adhesion molecule (ICAM)-1, vascular cell adhesion molecule (VCAM)-1, etc.), and a reduction in the endothelial production of nitric oxide (NO) which, in contrast to the other molecules quoted above, increases in production when the shear stress is laminar.^{48–50}

The specificity of atherosclerosis versus other forms of inflammation is essentially exerted at the endothelial level, where the early selective expression of adhesion molecules for monocytes allows these specific cells—and not neutrophils and lymphocytes—to adhere preferentially to the arterial endothelium, contrary to what occurs in other forms of inflammation. The immunoglobulin VCAM-1 is, in this context, very relevant in so far, through the interaction with the integrin very late activation antigen-4 (VLA4), expressed on monocytes and some lymphocyte classes.^{51,52} VCAM-1-VLA4 binding determines the selective recruitment of this particular type of leukocytes. The expression of VCAM-1 by the endothelium is favored, besides by a high laminar shear stress and turbulent flow, by virtually all cardiovascular risk factors. As examples, stimuli to its expression are modified or oxidized LDL, advanced glycation endproducts found abundantly in diabetes and uremia, primary inflammatory cytokines such as interleukin (IL)-1 and tumor necrosis factor (TNF)- α , as well as bacterial lipopolysaccharide (LPS).^{53,54} These substances are therefore considered the soluble mediators of vascular atherosclerotic disease, which conjoin with biomechanical factors in determining the extent and localization of atherosclerosis.

As a confirmation of the relationship of inflammatory phenomena with atherosclerosis, one may also remark the associations of atherosclerosis with classically “inflammatory” diseases. Rheumatoid arthritis, as an example, is associated with a high incidence of coronary heart disease. Both diseases show similarities in the activation of T-lymphocytes and mast cells, as well as in the production of TNF- α , IL-6, and adhesion molecules. The production of these and other cytokines by the synovial tissue in subjects with rheumatoid arthritis has similari-

ties with the proatherogenic phenomena occurring in subjects with vascular disease, such as conditions of insulin resistance, dyslipidemias, endothelial activation, and a prothrombotic phenotype. Patients with rheumatoid arthritis feature high levels of C-reactive protein (CRP), a marker of inflammation associated with increased cardiovascular risk.⁵⁵

The endothelium reacts to the proatherogenic activation with the production of counter-regulatory factors such as prostacyclin (PGI₂), NO, heparin-like complex polysaccharides able to activate antithrombin, type 1 plasminogen activator (PAI-1), protein S, thrombomodulin, and the secretion of ILs with antiproliferative and anti-inflammatory properties. In subjects with multiple risk factors for atherosclerosis, a simultaneous reduction in the endothelial production of such athero- and thrombo-protective mechanisms occurs.

The changes in the normal homeostatic balance between athero-protective and proatherogenic factors is generally explained by the activation, under the pressure of atherogenic risk factors, of a common proinflammatory mechanism, which in turn orchestrates the “pleiotropic” appearance of the endothelial proatherogenic phenotype. Such common mechanism may mostly be the transcription factor nuclear factor- κ B (NF- κ B) and a few other transcription factors. These represent the “central hub” of inflammation, a single intracellular system on which proinflammatory/proatherogenic stimuli converge determining its activation; and from which the various aspects of the pleiotropic endothelial proinflammatory phenotypes branch out (Fig. 1). Because of the multiplicity of the activation pathways (whereby the lack of one single activating stimulus may be compensated by the higher level of activation of other factors: the “redundancy” of the system), it is difficult to imagine that limiting the actions of one single risk factor may eradicate vascular disease, but it is logical to conceive that it can only serve to limit the problem. As a matter of fact, this intuitive approach to the choice of therapeutic targets in vascular disease has yielded appreciable results as the main current therapeutic strategies in vascular disease. One important consequence of this reasoning is also however that the central hub of inflammation (NF- κ B, with the possible accessory contribution of other transcription factors, such as AP-1, GATA, egr-1), may be a reasonable target in vascular inflammation.⁵⁶

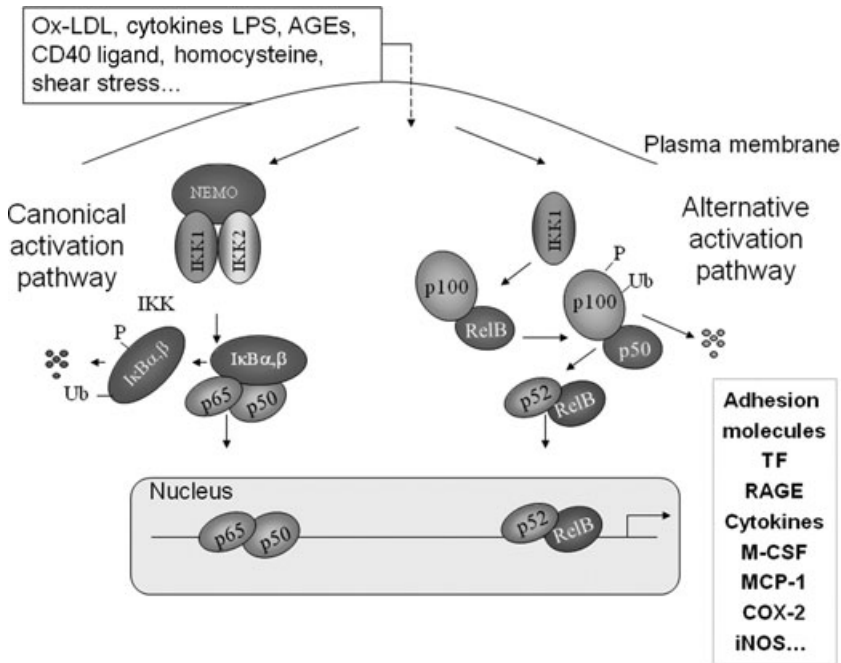


Figure 1. Canonical and noncanonical pathways of activation of the transcription factor nuclear factor (NF)- κ B. The NF- κ B system comprises a series of heterodimeric molecules (monomers including Rel-A/p65, p50, p52, Rel-B, c-Rel) normally sequestered in the cytoplasm and bound to an inhibitor (I- κ B). Under the influence of primary activating triggers (upper left inset), among which include inflammatory cytokines, modified LDL, etc., the increased generation of reactive oxygen species leads to the binding of I- κ B with ubiquitin and the proteolytic degradation of I- κ B. The heterodimers, made free from the inhibitor, can then migrate into the nucleus, where they bind specific recognition (consensus) sequences in the promoter region of a variety of genes of endothelial activation, with the final result of increased transcription of the respective genes (right lower inset). The partial overlap in properties of the primary triggers (“redundancy,” e.g., the nearly identical pattern of endothelial activation evoked by IL-1 and TNF), results in a similar cellular activation by different stimuli. The fact that multiple genes, e.g., E-selectin, VCAM-1, and ICAM-1, are expressed even when a single cytokine activates the system epitomizes the “pleiotropy” of the system, another property of inflammatory cytokines. Reproduced, with changes, from De Caterina, R., A. Zampolli, G. Lazzerini & P. Libby, 2007. Endothelial activation and the initiation of atherosclerosis. In *Endothelial Dysfunctions in Vascular Disease*. De Caterina, R. & P. Libby, Eds.: 26–35. Blackwell-Futura. New York, with permission by the publisher.

New biomarkers

This new knowledge on the central role of inflammation in atherogenesis has indeed yielded an entire new series of potential plasma biomarkers, different from those commonly used previously (e.g., based on the lipid profile), which largely overlap in reporting on pathogenetic mechanisms in vascular disease. Potential biomarkers here include VCAM-1, TNF- α , IL-1, IL-18, MMPs, CD40L, adiponectin, PAI-1, fibrinogen, and several others.⁴⁰ The ideal biomarker should be easily measured, have minimum circadian variations, and have a long half-life. CRP closely matches such a description.⁵⁷ Conversely, as an example, PAI-1 has a half-life of about 6 h and features important circadian variations.

CRP is predictive of cardiovascular events (myocardial infarction, stroke, and sudden cardiac death) in addition to measurements of LDL cholesterol.⁵⁸ For this reason, its measurement has been proposed in addition to the list of classical risk factors based on the Framingham algorithm.⁵⁹ Subjects with CRP < 1 mg/L are at low risk, whereas subjects with levels > 3 mg/L are at high risk. Risk of ischemic heart disease and stroke in the latter situation indeed is increased by a factor of 1.3 and 1.6, respectively).⁶⁰ Conversely, the evidence that CRP may also be a mediator of vascular disease, and therefore itself a potential therapeutic target, appears much weaker, especially on the basis of recent Mendelian randomization studies.⁶⁰ Other new potential biomarkers stemming out of our recent

knowledge on vascular inflammation include CD40 ligand,⁶¹ adiponectin (as an inverse correlate),⁶² IL-18,⁶³ and some MMPs, potentially reporting on the risk of plaque rupture,⁶⁴ as well as TF, marking a prothrombotic phenotype.⁶⁵

New therapeutic approaches

A hypothesis remains a hypothesis, no matter how attractive it is, until one can fully demonstrate its validity. The demonstration of the causality relationship for the inflammatory hypothesis of atherosclerosis requires the testing with adequate therapies. We have several therapies available with an impact on vascular atherosclerosis, but only few of them are practicable on a large scale in prevention strategies. These may largely be classified in three main groups:

1. therapies acting on primary atherogenic triggers,
2. therapies acting on the final effectors, and
3. therapies acting on the central common pathway of vascular inflammation.

We will briefly discuss these and highlight where novelties in this area may appear.

Therapies acting on primary atherogenic triggers

To combat inflammation linked to atherosclerosis, a first logical approach, followed for decades, has been to limit the classical risk factors of vascular diseases, with interventions on diet, physical activity, the cessation on smoking, or the effects on specific risk factors. Changes in lifestyles are unfortunately a constant source of frustration. The latest EuroAspire survey, which has produced a snapshot of practices of secondary prevention in eight European countries in 2005, comparing it to previous surveys of the same kind carried over in 1996 and 1998,⁶⁶ has shown how far we are from reaching the declared objectives. We are much better in prescribing drugs, mainly antihypertensive drugs and statins, than to induce people to stop smoking, reduce caloric intake or adopt healthy lifestyles. In any case, we are doing something good for atherosclerosis when we limit risk factors such as LDL cholesterol (with statins) or hypertension (with antihypertensive drugs).

In doing all this, we also reduce vascular inflammation. Statins are a particularly interesting class of drugs in this respect. Beyond reducing LDL chole-

sterol, statins also reduce markers of vascular inflammation, including IL-6, TNF, and CRP.^{67,68} If a cholesterol-lowering therapy is anti-inflammatory, it should reduce CRP. This is exactly what happens: CRP is reduced by 15–50% with statins,^{57,69–72} in what appears to be a class effect. If a reduced CRP really is a measure of the probability of cardiovascular complications, cardiovascular risk should parallel such a reduction. This is exactly what happens. A recent study has indeed reported that in patients with an acute coronary syndrome receiving statin therapy, subjects with CRP < 2 mg/dL had a reduced risk of reinfarction or vascular death compared with subjects with higher values (2.8 vs. 3.9 events per 100 persons-year; $P < 0.006$).⁵⁸ This benefit appears somehow independent from reduction in LDL cholesterol. Therefore, the clinical benefit of statins seems to parallel anti-inflammatory effects. In any case, however, LDL reduction is itself an anti-inflammatory therapy, and this activity appears central to the beneficial effect of statins.

A similar reasoning can be done for drugs inhibiting the renin–angiotensin system, angiotensin-converting enzyme (ACE) inhibitors, and angiotensin receptor blockers, commonly used throughout the spectrum of vascular disease.^{73,74}

Therapies acting on the final effectors

Classical anti-inflammatory therapies are by many considered the essential ingredients of a therapeutic cocktail destined to demonstrate causality of inflammation in atherosclerosis. However, many classical anti-inflammatory therapies have contraindications in patients with vascular disease. Glucocorticoids, for example, were experimented with in the 1970s in the acute phase of a myocardial infarction, but their development with this indication was quickly halted owing to a higher incidence of heart rupture. Traditional nonsteroidal anti-inflammatory drugs, as well as more recently developed cyclooxygenase (COX)-2 selective inhibitors (coxibs), have a theoretical contraindication in that they limit vascular production of prostacyclin, which is mainly COX-2-dependent, and may compete for the antithrombotic effects of aspirin. Anticytokine drugs, which selectively hit one of the many final targets of inflammation, have been tested serendipitously. A therapy with etanercept, a derivative of a monoclonal antibody against TNF- α , has been experimented in a phase 3 study in heart failure (the RENEWAL study),

without appreciable results even limited to purely ischemic endpoints.⁷⁵ The MRC-ILA-HEART trial plans to study 186 patients with non-ST elevation myocardial infarction in the United Kingdom using anakinra, and IL-1 receptor antagonist (IL-1Ra). The study will evaluate efficacy and safety of the drug on circulating levels of CRP, as a prelude to a possible efficacy phase 3 study.⁷⁶ Similar approaches are the possible interferences with the production or activity of leukotrienes, with action at the level of 5-lipoxygenase or at leukotriene receptors. Although conceptually interesting, such approaches interfere with only some of the multiple mediators of inflammation potentially involved.

The case of dipyridamole

The total suppression of NF- κ B activity, when tried in animal experiments, interferes with embryogenesis and with immune functions, and does not therefore appears as a viable option. On the contrary, first in *in vitro* and then *in vivo* in animal experiments, several drugs generically defined as “antioxidant” have been tested. Such drugs interfere with the generation of some reactive oxygen species, which are in turn activators of NF- κ B. One such substance, succinobucol, has been brought to phase 3 in non-ST elevation acute coronary syndromes within the ARISE study, with mostly disappointing results.⁷⁷ One should however mention that the extent of inhibition of NF- κ B with the *in vivo* administration of this drug was not as well documented as *in vitro*. Methotrexate, a drug commonly used in rheumatic diseases and relatively safe at low doses, might be an alternative worth testing, and this approach is currently being pursued.

An alternative option in this respect is to revisit old drugs for which some clinical efficacy has been already documented, and try to exploit novel notions about their mechanism of action. One such drug is dipyridamole. Originally branded as an antiplatelet agent, the drug has still an uncertain role in coronary heart disease (due to lack of data from well-performed clinical trials), but has evidence of efficacy in the prevention of ischemic stroke.⁷⁸ In our own hands (unpublished data), the drug reduces monocytoïd cell and endothelial cell pro-MMP9 and MMP-9 after TNF α /phorbol myristate acetate or high-glucose stimulations, thus potentially reducing mechanisms of instabilization of atherosclerotic plaques (for the effect on monocytes, the main

secretors of collagenolytic enzymes in the plaque, and for the effect on endothelial cells, where reduction on MMPs should lead to reduced plaque angiogenesis). Such effects appear to occur through inhibition of NF- κ B activation. At the endothelial level, notably, dipyridamole induces endothelial cell COX-2 expression, alone or after TNF α /PMA stimulation, therefore not potentially curtailing the production of endothelial prostacyclin. These observations are intriguing, although their clinical relevance remains to be demonstrated.

Conclusions

Inflammation plays a central role in vascular disease, from plaque inception to plaque progression and instabilization. The now demonstrated sequence of events linking inflammation with vascular disease differs substantially from the old idea of vascular disease as a problem of lipid accumulation. Inflammatory markers such as CRP, derived from such knowledge, merit careful consideration to be included in our algorithms of risk evaluation. Knowledge of the mechanisms of vascular inflammation also holds the promise of unraveling new targets for therapeutics. Such new therapies may include the revisitation of the mechanism of action of old drugs.

Conflicts of interest

The authors declare no conflicts of interest.

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