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TOPIC HIGHLIGHT

Possible Pivotal Role of Latent Chronic *T. gondii* Infection in the Pathogenesis of Atherosclerosis

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ABSTRACT

Cardiovascular diseases such as atherosclerosis, atherothrombosis, coronary artery disease and stroke are the main causes of death worldwide. Atherosclerosis is a chronic, progressive immune-inflammatory and fibroproliferative disease of medium and large sized arteries with increased blood lipoprotein/cholesterol and their disposition in the arterial wall, important T_H1 type proinflammatory reaction, and thrombogenic status. Elevated plasma levels of proinflammatory cytokines have been demonstrated in patients with established atherosclerotic disease, and therefore it is believed that cytokines are key players in all stages of disease and have a profound influence on the pathogenesis of the disease. Several authors suggested that the number of different pathogens, including *T. gondii*, may promote synergistic inflammatory responses that are capable of triggering and exacerbating atherosclerotic process. Latent chronic

T. gondii infection may be a frequent cause of vascular endothelial cell dysfunction because this pathogen attacks all nucleated cells, and endothelium cells have enhanced susceptibility to infection with T. gondii tachyzoites. Oxidative stress characteristic for atherosclerosis may be caused by the host infection with the parasite. T. gondii is unable to synthesize sterol and acquires cholesterol from the host LDL receptor pathway. The accumulation of foamy transformed macrophages in the aortic intima characteristic for atherosclerotic lesion is due to acyl-CoA: cholesterol acyltransferase 1 and 2 (ACAT1 and ACAT2) increased expression in macrophages and other cells infected with T. gondii. ACAT2 is crucial in foam cell formation and development of atherosclerosis. The parasite also expresses two cholesteryl ester(CE)-synthesizing enzymes TgACAT1 and TgACAT2 that contribute to the CEs formation for storage in lipid bodies, but at the same time ACAT and CEs play a crucial role in replication of the pathogen. It must be emphasized that the increased expression of ACAT1 and ACAT2 normally present in macrophages may thus be further enhanced by the superimposing TgACAT1 and TgACAT2 activities available in the host cells infected with T. gondii. In addition, proinflammatory cytokines increase foam cell formation and latent chronic T. gondii infection persistently generates these biomediators and thus play an important role in foam cell biogenesis. Patients with atherosclerosis had increased plasma levels of TGF-β and this cytokine increased T. gondii replication in the host cells, and participated in development of fibrotic changes in atherosclerotic lesions. The parasite may also produce some extracellular vesicles known to play a role in triggering cardiovascular events. Atherogenesis involved platelets activation with subsequent serotonin release. Hyperserotoninemia was also reported in autistic and mentally retarded children, and recently, a significantly higher seroprevalence of chronic toxoplasmosis was found in autistic children as compared with controls. T. gondii infection caused also increased leptin levels, and a marked association between T. gondii seropositivity and obesity has been demonstrated. Cysteine cathepsins (Cat) also play an important role of in foam cell formation and generation of amyloid in atherosclerotic arteries. Infection of vascular endothelial cells with T. gondii tachyzoites provide an additional source of Cat that are superimposing on similar enzymes already present in host cells, because the parasite expresses few members of the cathepsins, such as CatL-like, CatB-like, and CatClike proteases. Finally, vitamin D exerted beneficial effects in both atherosclerosis and T. gondii infection, especially that it improved host immunity and decreased proliferation of T. gondii tachyzoites in

macrophages and reduced tissue pathology caused by the pathogen. Because the parasite is widely disseminated in animals and humans, it seems that latent chronic toxoplasmosis play a critical role in the pathogenesis of atherosclerosis.

Key words: Atherosclerosis; Atherothrombosis; Cardiovascular diseases; Stroke; *T. gondii* Infection; Endothelial dysfunction; Oxidative stress; ACAT1; ACAT2; Foam cells; Mast cells; Blood coagulation; Iron; Leptin; Extracellular vesicles; Platelets; Serotonin; Cysteine cathepsins; Vitamin D; Statins; Propranolol

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INTRODUCTION

Atherosclerosis

Cardiovascular diseases (CVDs) are the leading cause of death worldwide, accounting approximately 17.3 million deaths per year^[1,2], and this number will probably increase to 23.3 million by 2030^[3]. Atherosclerosis also is the main cause of death, and in Europe, cardiovascular disease, atherothrombosis, coronary artery disease and stroke account for over 4 million deaths per year^[4].

Atherosclerosis is a chronic, progressive immune-inflammatory and fibroproliferative disease of medium and large sized arteries with three main components: (a) increased blood lipoprotein/cholesterol and their disposition in the arterial wall; (b) an important T_H1 type proinflammatory reaction; and (c) thrombogenic status^[5]. The development of atheromatous plaques involves a number of cellular and molecular events including endothelial dysfunction, leukocyte recruitment and diapedesis, low-density lipoprotein (LDL) transcytosis and oxidation, intimal smooth muscle cells recruitment and proliferation, synthesis of extracellular matrix proteins, and accumulation of foamy macrophages along with interplay of various cytokines^[6].

Atherosclerotic process is initiated by the infiltration, retention and accumulation of LDL in the $\it tunica$ intima of the arterial wall[7-9], and typically begins with endothelial cell dysfunction, mononuclear cell recruitment into vascular wall, and the differentiation of macrophages into foam cells[10,11]. Atherosclerotic lesions are characterized by progressive accumulation of lipids, macrophages, NK cells, T and B lymphocytes, smooth muscle cells, and fibroproliferative elements in the $\it intima$ of arteries[12,13]. Functional analysis of different T-cell subsets role identified the $T_{\rm H}1$ responses as proatherogenic, whereas $T_{\rm H}2$ immune responses exert antiatherogenic actions[14,15]. Atherosclerosis is also associated with B-cell activation, and traditional T cell-driven B2 cells responses promote atherosclerosis, while innate B1 cells provide protective action through the secretion of naturally occurring antibodies[14,15].

Several authors have hypothesized that the number of different pathogens, including T. $gondii^{[16]}$, may promote synergistic inflammatory responses that are capable of triggering and exacerbating atherosclerosis^[17]. Lipopolysaccharide was found to be a potential contributor to the development of atherosclerotic plaque even at extremely low serum concentrations^[18,19]. In susceptible individuals, the presence of a chronic infective stimulus, as well as a prolific generation of proinflammatory cytokines, may directly

activate metabolic pathways, such as for example cholesterol esterification, leading to foam cell formation and development of atherosclerosis^[20]. Macrophages and foam cells present in the evolving atherosclerotic lesion, secrete various proinflammatory cytokines, such as TNF-α, IFN-γ, IL-6, and chemokines, including ICAM-1 and VCAM-1. In response to enhanced production of proinflammatory cytokines, endothelial cells express high levels of leukocyte adhesion molecules on their surface, leading to further mononuclear cell recruitment and, hence, to development of a chronic inflammatory state^[21-23].

T. gondii infection

Toxoplasmosis is one of the most frequent infections in both animals and humans affecting approximately 30-50% of the total world human population^[24,25]. Ophthalmoimmunologists suggested that even some six billion people are chronically infected with *T. gondii*, with high seroprevalence in tropical areas and low prevalence in cold regions^[26]. Seroprevalence increased markedly with age, to approximately 80-100% in persons aged 66-75 years^[26,27]. At present, however, in immunocompetent individuals *T. gondii* infection is still believed to be asymptomatic^[28,29], but a rapidly increasing body of literature strongly indicate that the parasite is slowly emerging as a global health threat^[24,25,30].

Three clonal strain types (I, II, and III) of T. gondii predominate worldwide^[31]. In mice, the outcome of infection is highly dependent of the parasite genotype with type I strains being uniformly virulent, while types II and III are nonvirulent. Strains types II and III result in controlled infections that persist chronically^[32]. The most prevalent in nature and frequent in humans infection with type II strain parasites stimulate NF- κ B nuclear translocation and lead to an abnormal overproduction of proinflammatory $T_H 1$ cytokines, including IFN- γ , TNF- α , IL-1 β , IL-12 and IL-6, that are typically protective^[31-34]. Different strains of T. gondii produce different cytokines and cause different clinical entities^[30], e.g. type I strain is more prevalent in ocular toxoplasmosis^[30,35].

The aim of the work was to review and critically analyze available literature data, especially focusing on the molecular pathomechanisms of both atherosclerosis and toxoplasmosis, and provide solid scientific documentation that latent chronic *T. gondii* infection probably play an essential role in triggering, development and progression of atherosclerosis in both animals and humans.

INCREASED SUSCEPTIBILITY OF VASCULAR ENDOTHELIAL CELLS TO INFECTION WITH T. gondii MAY BE A FREQUENT CAUSE OF THE ENDOTHELIAL DYSFUNCTION IN PATIENTS WITH ATHEROSCLEROSIS

Endothelial cells maintain the functional integrity of the vascular wall, and are involved in controling vascular wall permeability, the regulation of immune and inflammatory reactions, the modulation of blood flow and vascular resistance, and the maintenance of a non-thrombogenic blood-tissue interface^[36].

Damage of vascular endothelial cells by proatherosclerotic risk factors, such as proinflammatory cytokines, oxidized proteins/lipids, and/or hyperglycemia, leads to endothelial dysfunction^[37,38]. Accelerated apoptosis of vascular endothelium cells^[39] and an associated delay in cell cycle progression, affecting morphology and impairing function of the endothelial monolayer, are believed to be key bioevents for triggering and progression of micro- and

macrovascular diseases, including premature atherosclerosis and diabetic neuropathy [37-39].

Endothelial cell dysfunction plays an important role in the process of atherosclerotic lesion formation by promoting leukocyte adhesion, increasing chemokine generation and cell permeability to lipids, enhancing LDL oxidation, stimulation of vascular smooth muscle cell proliferation and migration, and platelet activation^[40]. Endothelial dysfunction is characterized by a reduction in the bioavailability of vasodilators, especially NO, and the increase in production of endothelium-derived contractile factors. There is a state of endothelial cell activation associated with increased generation of proinflammatory, proliferative, and procoagulant biofactors, all favoring triggering and development of atherogenesis^[41]. Endothelial dysfunction may be caused by acute and/or chronic inflammation state and enhanced production of proinflammatory cytokines and free radicals ROS and RNI. Increased LDL oxidation causes enhanced expression of adhesion molecules in the endothelium, which facilitates monocyte infiltration of subendothelial space^[42].

It must be noted that flow-mediated dilation (FMD) (the most widely used non-invasive ultrasound method to assess endothelial function) derived using a proximal or distal cuff may serve as future cardiovascular events prediction^[42a,42b]. Age- and sex-specific FMD data were early markers of atherosclerosis, and males had a lower FMD values than females, and FMD appeared to decline with age (except of pediatric population)^[42c-42e]. Obstructive sleep apnea and hypertension also appeared to be early clinical markers of carotid atherosclerosis^[42f,42g]. In addition, plasma pentraxin 3 (PTX3) was found to be a more sensitive predictor of endothelial dysfunction than serum high-sensitive C-reactive protein (hsCRP)^[42h], at least in part because PTX3 overexpression in neutrophils was associated with intracellular ROS overproduction^[42i].

Vascular endothelial cells are susceptible to infection with *T. gondii* tachyzoites

The parasite is disseminating in the body as a Trojan horse in various eukaryotic cells, including endothelial cells and macrophages, and division rate of intracellular unprimed *T. gondii* tachyzoites in endothelial cells, monocyte-derived macrophages, peritoneal, or alveolar cells is rapid (**Table 1**)^[43].

In congenital toxoplasmosis, the parasites first invaded the umbilical vein endothelial cells and were then disseminated throughout the fetus^[60,61]. Canedo-Solares *et al*^[60] found that RH tachyzoites infected up to 19% human umbilical vein endothelial cells (HUVECs), while ME49 parasites invaded up to 63% human microvascular endotheial cell-1 line (HMEC-1). Both RH and Me49 tachyzoites invaded 46 and 49% HUVECs and HMEC-1 cells, respectively. Reinvasion and formation of new parasitophorous vacuoles of infected cells was more frequent than invasion of noninfected cells. Thus, dissemination and vertical transmission of *T. gondii* appeared to be parasite strain type, host cell type/subtype and activation state- dependent. Replication speed of the parasite seemed to affect its virulence^[60].

Smith *et al*^[62] found that retinal vascular endothelium cells have enhanced susceptibility to infection with *T. gondii* tachyzoites in comparison with aorta (55% more), umbilical vein (33%), and dermal endothelial cells (34% more), and tropism of different pathogens for particular cell types and/or specific tissue sites was a long-recognized biological phenomenon^[62]. Free *T. gondii* tachyzoites had the ability to transmigrate a stimulated human retinal endothelium monolayer^[63]. Tachyzoites crossed retinal endothelium assisted by intercellular adhesion molecule-1 (ICAM-1) (the cell surface IgG immunoglobin

Table 1 Division rate of intracellular T. gondii tachyzoites in primary human cells in vitro (acc. to Channon $et \, all^{[43]}$; with own modification).

Cell type	Parasite division rate Unprimed	IFN-γ primed	Mechanism	Refs
Hemopoietic				
Lymphocyte	S			[43]
Neutrophil	S			[43-45]
Adherent monocyte	S			[45-50]
Nonadherent monocyte	R	R	ROS; not TS	[43,47]
Dendritic cell	R			[43]
Alveolar macrophage	R	S	Partly TS	[50]
Peritoneal macrophage	R	S		[51]
Monocyte-derived macrophage	R	S	ROS; not RNI	[49-52]
Nonhemopoietic				
Neuron	S			[53]
Foreskin fibroblast	R	S	TS	[54,55]
Umbilical vein endothelial cell	R	S	TS or ROS	[49,56]
Retinal pigment epithelial cell	R	S	TS	[57]
Fetal astrocyte	R	S	RNI	[53,58]
Fetal microglial cell	R	R		[59]

R, rapid; S, slow; RNI, reactive nitrogen intermediates; ROS, reactive oxygen species; TS, tryptophan starvation.

superfamily member) in vitro, and ICAM-1 blockade significantly (by approximately 50%, p < 0.001) inhibited the parasite migration across stimulated human retinal, but not choroidal, vascular endothelium^[63]. There has been interest in the heterogeneity of vascular endothelium, not only between arteries and venous types^[64], but also between same type vessels located within different organs or within different tissues in the same organ^[65,66]. An intracellular environment that is low in ROS or RNI and/or high in iron or tryptophan is known to facilitate tachyzoite proliferation^[49,54,67]. Courret et al^[68] tracked fluorescently labeled tachyzoite-infected CD11c-positive or CD11b-positive leukocytes from blood to brain, and showed that human dendritic cells (DCs) and monocytes were more permissive to infection with T. gondii tachyzoites than neutrophils or lymphocytes^[43]. Moreover, transmigration of human DCs across retinal endothelium was augmented following infection with the parasite^[69]. In this context, it must be emphasized that nearly 1 million strokes occur in the US each year, and approximately 10-15% can be attributed to intracranial atherosclerotic disease^[70], and the infection of brain vascular endothelial cells with tachyzoites may play an important role in these clinical bioevents.

Enhanced T. gondii adhesion to vascular endothelium during disturbed blood flow

Atherosclerosis disease occurs at sites of disturbed blood flow, typically in the curvature of vessels or at branching points^[71-76]. Smooth laminar blood flow induces NO production, which has multiple anti-atherosclerotic functions, including suppressing adhesion molecule expression by endothelial cells, inhibiting platelet aggregation, and inhibition proinflammatory cytokine production^[77]. It was demonstrated that *T. gondii* modulated the dynamics of human monocyte adhesion to vascular endothelium under fluidic shear stress^[78]. *T. gondii*-infected primary human monocytes and THP-1 cells exhibited altered adhesion dynamics compared with uninfected monocytes. Infected cells rolled at markedly higher velocities (2.5- to 4.6-fold) and over greater distances (2.6- to 4.8-fold), than

uninfected monocytes, before firmly adhering^[78]. Moreover, *T. gondii* tachyzoites adhered and glided on human vascular endothelium under shear stress conditions^[79]. Compared to static conditions, shear stress enhanced *T. gondii* helical gliding, resulting in a significantly greater displacement, and increased the percentage of tachyzoites that invaded and sustained adhesion to vascular endothelium. It appeared that the parasite surface protein MIC2 contributed to initial adhesion but was not required for adhesion strengthening^[79].

Increasing T. gondii infection percentage during human aging

Endothelial cells are a replicative niche for entry of the parasite to the brain vasculature, and replication in and lysis of these cells precedes invasion of the central nervous system^[80]. Vascular endothelial dysfunction occurs during the human aging process and is accompanied by deterioration in the balance between vasodilator and vasoconstriction substances produced by the endothelium^[81]. This imbalance is mainly characterized by a progressive reduction of the bioavailability of NO and an increase in the production of COX-derived vasoconstrictor factors. Both circumstances are in turn related to an increased generation of ROS and RNI^[81]. Interestingly, the percentage of individuals chronically infected with *T. gondii* also is increasing with age (**Table 2**) and this may, at least in part, be responsible for development of vascular endothelial dysfunction during aging.

Proinflammatory and antiinflammatory cytokines acting as positive or inhibitory stimuli influence the development and maintenance of differentiated properties in the vasculature. In SMCs, the gene for the contractile protein, α -actin, is stimulated by TGF- β but inhibited by IFN- $\gamma^{[82,83]}$. Similarly, TGF- β strongly promoted the synthesis of interstitial collagens types I and III by human SMCs, whereas IFN- γ powerfully inhibited their synthesis of collagen, as well as α -actin^{[84]}. Vascular cells participate in and propagate the inflammatory response at sites of microbial challenge or pathological processes by generating large amounts of cytokines (**Table 3**). Both ECs and SMCs responded to stimulation with IL-1, TNF- α , or CD40 ligand by producing great quantities of IL-6[85], and in one study of 14,916 apparently healthy men who developed myocardial infarction, plasma IL-6 levels were significantly elevated as compared with control individuals who did not have heart problems^[86].

Impaired nitric oxide (NO) production in endothelial dysfunction

Vascular endothelium-derived NO, a major mediator of endotheliumdependent vasodilation, has important antiinflammatory and antithrombotic properties, such as inhibiting leukocyte adhesion, limiting platelet adhesion and aggregation, and the expression of PAI-1, a prothrombotic protein[88-90]. A common mechanism underlying endothelial dysfunction relates to inflammation and increased vascular production of ROS and RNI^[90]. Endothelial cell tetrahydrobiopterin (BH₄), a naturally occurring essential cofactor for several critical metabolic pathways, including the production of NO, is required for the maintenance of vascular function through enhanced NO production, and reduced levels of BH4 were associated with vascular dysfunction^[76]. Stroes et al^[88] demonstrated that BH₄ restored vascular endothelial function in patients with familial hypercholesterolemia which was associated with impaired NO activity. Recently, it was also however reported that although acute and short-term BH₄ oral administration improved vascular endothelial function in patients with rheumatoid arthritis, the supplementation did not alter aortic stiffness (an independent risk factor for CVD[89]. The authors suggested that these two disturbances may occur in parallel and probably share common risk factor such as inflammation.

Table 2 Percentage of *T. gondii* positive individuals among 214 nonpsychiatrically affected controls depending on age analyzed during a large epidemiologic study of 869 psychiatric patients (Hinze-Selch *et al*^[27]).

Percent						Age	(yrs)					
T. gondii	18-	21-	26-	31-	36-	41-	46-	51-	56-	61-	66-	71-
positive	20	25	30	35	40	45	50	55	60	65	70	75
100												100
80									80		80	
60							58	58		60		
40		35	32	40	39	38						
20	19											
0												

In the control individuals 45 yrs or younger recruited from the same geographical region as the psychiatric patients admitted to the hospital, serofrequency of T. gondii infection ranged between 20 and 40% without any systematic age effect, whereas in the individuals older than 45 yrs serofrequency systematically increased with age from about 40% to almost $100\%^{[27]}$.

Table 3 Regulation of human vascular endothelial and smooth muscle cells function by positive and inhibitory stimuli (acc. to Hansson *et al*^[82]; with own modification).

Target	Positive stimuli	Inhibitory stimuli
SMC proliferation	PDGF (TGF-β)	IFN-γ (TGF-β)
SMC matrix/collagen	TGF-β (IL-1, PDGF)	IFN-γ
SMC contractile proteins	TGF-β	IFN-γ
Ig-type adhesion molecules	IL-1, TNF-α, IFN-γ, CD40L	
SMC NOS2	IL-1, TNF-α, IFN-γ	TGF-β, IL-4
EC/SMC COX-2	IL-1, TNF-α, IFN-γ, CD40L	
EC E-selectin	IL-1, TNF-α, IFN-γ, CD40L	
EC/SMC tissue factor	IL-1, TNF-α, CD40L	
EC/SMC MMPs	IL-1, TNF-α, CD40L	IFN-γ

EC, endothelial cell; Ig indicates immunoglobulin superfamily; CD40L, CD40 ligand (functional CD40L is expressed on human vascular ECs, SMCs, and macrophages; the activated T cells also express CD40L and in advanced human atherosclerotic plaques T cells constitute 10-20% of the cell population^[53,55]); COX-2, cyclooxygenase 2; NOS2, nitric oxide synthase 2; MMPs, matrix metalloproteinases; PDGF, platelet-derived growth factor; SMC, smooth muscle cell.

Table 4 Intracellular pteridine levels in cultured and freshly isolated human umbilical vein endothelial cells (HUVEC) (acc. to Rosenkranz-Weiss $et~all^{[91]}$;with own modification).

	Pteridine levels				
HUVEC conditions	Biopterin	Neopterin			
	pM per million cells				
Cultured, passage 3-6					
Control (4)	< 2	< 2			
TNF, IL-1β, and IFN-γ (4)	27.6 ± 6.7 °	24.6 ± 11 a			
Cultured, primary (3)	< 2	< 2			
Freshly isolated (3)	31.7 ± 1.3 ^a	ND			

Results are mean \pm SEM. ^a p < 0.05 vs control cultured HUVEC. Numbers in parentheses represent the number of samples. ND, not done.

Rosenkranz-Weiss *et al*^[91] demonstrated that proinflammatory cytokines such as IFN- γ plus TNF or IL-1 β enhanced NOS specific activity in cultured human endothelial cells by increasing intracellular BH4 levels (**Table 4**).

Effect of other putative vascular cell pathogens on endothelial function

The first line of immune host defense is based on detection of pathogen-associated molecular patterns that evoke inflammatory and toxic response^[92,93]. Several studies suggested a role for *Chlamydia pneumoniae* (CP) in atherosclerosis^[21]. For example, HSP60 of this microbe resembles human HSP60 and can elicit inflammatory response^[94]. Chlamydial and human HSP60 induced TNF-α and MMP production by macrophages^[95] and stimulated E-selectin, ICAM-1, and VCAM-1 expression on endothelial cells^[94]. HSP60 also markedly enhanced IL-6 production by endothelial and smooth muscle cells, and macrophages^[94]. Another putative vascular pathogen, cytomegalovirus (CMV), encoded a chemokine receptor that rendered infected smooth muscle cells susceptible to CC chemokine-induced migration^[96].

Several authors demonstrated the relationship between the seropositivity to CP, HSV-1 and 2, HCMV, HAV, HP, EBV, and atherosclerosis/coronary artery disease[97-101]. Recently, it was also suggested that more infectious agents rather than any single pathogen may be involved in the development of atherosclerosis and the subsequent cardiovascular events because the proatherogenic effect of each infectious organism might be amplified^[1,102]. Specifically, in 86 patients with no angiographically significant CAD, Prasad et $al^{[103]}$ found a significant correlation between total infection burden to five investigated agents (CP, CMV, H. pylori, hepatitis A, and herpes simplex virus) and decreased intracoronary endotheliumdependent vasodilation (p = 0.03). [Of note, the eventual presence of latent chronic T. gondii infection in these patients had not been investigated]. The inflammatory state can induce oxidative stress by enhancing the production of ROS in the vascular wall, therefore contributing to the progression and destabilization of atherosclerotic plaque and consequently to CADs[104]. Key process in the development of atherosclerosis is LDL oxidation and accumulation in vascular cells, promoting foam cell formation, as well as increased secretion of mediators of inflammation, including IL-1, IL-6 and TNF- $\alpha^{[104]}$.

On the other hand, Khairy *et al*^[105] found lack of association between chronic infection with CP, CMV, EBV, HP, or pathogen burden, and endothelial function in 65 male individuals, aged 20 to 45 yrs (mean age = 29.3 ± 5.5 yrs), with no risk or known coronary artery disease(CAD), suggesting that these agents were not implicated as early etiologic triggers in the pathophysiology of CAD (**Table 5**). Endothelial function was determined by flow-mediated brachartery vasodilation and seroprevalence values were 65.1%, 34.9%, 88.9%, and 14.3%, for CP, CMV, EBV, and HP, respectively^[105], consistent with the earlier values reported in young patient populations^[106,107], and the coexistence of these infections has been well described^[108].

In summary, latent chronic *T. gondii* infection may be a frequent cause of vascular endothelial cell dysfunction because the parasite attacks all eukaryotic cells and endothelium cells have enhanced susceptibility to infection with *T. gondii* tachyzoites with rapid division rate. Increase of vascular endothelial dysfunction occurs during the human aging process and it was found that the percentage of individuals chronically infected with *T. gondii* also is increasing with age. Human studies showed that the infection burden such as *H. pylori*, CMV, and EBV, decreased endothelium-dependent vasodilation, but eventual additional latent chronic *T. gondii* infection in these participants had not been investigated.

OXIDATIVE STRESS CHARACTERISTIC FOR ATHEROSCLEROSIS MAY BE CAUSED BY LATENT CHRONIC T. gondii INFECTION

Chronic inflammation (innate immunity-associated) may trigger

Table 5 Flow-mediated brachial artery vasodilation caused by chronic low-grade infection with *Chlamydia pneumoniae*, Ebstein-Barr virus, cytomegalovirus, and *H. pylori*, associated with decreased endothelial function in seropositive healthy young men (acc. to Khairy *et al* ^[105]; with own modification).

Infectious agent	FMD Seropositive	FMD Seronegative	Parameter estimate 95% CI	P
Chlamydia pneumoniae	9.8 ± 4.5	8.5 ± 4.5	0.280 (-2.053, 2.613)	0.8151
Ebstein-Barr virus	9.5 ± 4.5	8.1 ± 4.2	0.159 (-0.432, 0.750)	0.6006
Cytomegalovirus	9.0 ± 4.5	9.5 ± 4.5	-0.806 (-3.403, 1.789)	0.5452
H. pylori	13.6 ± 4.4	8.7 ± 4.1	2.888 (-0.354, 6.129)	0.0888

FMD, flow-mediated brachial dilation; CI, confidence interval.

initial events that can lead to atherosclerotic cardiovascular disease^[109]. Early stages of chronic inflammation and onset of atherosclerosis may be evident in embryonic life^[110], and be perpetuated by classic atherosclerosis environmental risk factors, including *T. gondii* infection. Lipid peroxidation and formation of oxLDL are pivotal processes in the development of atherosclerosis that represent a crucial proinflammatory stimulus^[111,112]. Atherosclerosis and autoimmune diseases have a number of pathogenic similarities^[113], and autoimmunity plays an essential role in pathophysiology of atherogenesis. Atherosclerosis triggers the production of autoantibodies against substances such oxLDL^[113].

Oxidation of LDL result in the formation of reactive aldehyde products such as malondialdehyde (MDA), 4-hydroxynonenal (4-HNE), or 1-palmitoyl-2-oxovaleroyl phosphatidylcholine[114]. MDAmodified LDL is commonly utilized as an indicator for oxLDL, because MDA is the most abundant aldehyde arising from lipid peroxidation. Recently, Matsuo et al[115] demonstrated that circulating MDA-LDL concentrations were associated with the presence of thin-cap fibroatheromas in patients with coronary artery disease^[115]. Bardag-Gorce et al[116] showed that 4-HNE inhibited the proteasome activity and that the proteasome was unable to degrade proteins heavily modified by 4-HNE. Ultrastructurally, 4-HNE-treated cells displayed extensive vacuole formation, pinocytic body formation, crescent-shaped phagophores, and multilamellar vesicles[117]. In addition, 7-ketocholesterol present in oxLDL, not only triggers oxidative damage and protein-4-HNE modification but also extensive vacuolization in SMCs[118]. OxLDL is also a potent chemoattractant and induces the secretion of macrophage-chemotactic protein-1 by endothelial cells[119-121].

Acute or persistent oxidative stress results in an intracellular increase of ROS that damage the lysosomal membrane. When oxidative injury overcomes the cellular defenses, cells probably die via apoptosis[117,122]. The production of ROS and RNI (peroxynitrite, i.e. ONOO, a potent oxidant and nitrating agent that can cause formation of several lipid oxidation products including MDA, lipid peroxides and lipid hydroxides, and oxysterols, as well as interaction iron and copper species[123-125] has been implicated in atherosclerosis principally as means of damaging LDL[126-128]. Oxidation of lipids present in LDL generate a broad range of reactive products such as the above-mentioned aldehydes that are capable of reacting with lysine residues[123]. Specifically, the extent of oxidation of linoleic acid measured in atherosclerotic lesion is approximately 20% and is similar to *in vitro* studies using copper-mediated LDL oxidation^[129]. Antibodies directed to specific oxidized lipid-protein adducts cross react with material in atherosclerosis, and titers of autoantibodies to oxLDL are positively correlated with the progression of atherosclerotic lesions in humans[130].

Latent chronic T. gondii infection

Several studies suggested that infectious pathogens, including *T. gondii*, *H. pylori*, *Chlamydia peumoniae*, and *Enterococcus fecalis*^[131-133] may be responsible for inducing oxidative stress and triggering primary atherosclerotic lesions in both animals and humans^[131,134,135]. *T. gondii* infection may be at least partly responsible for development of these abnormalities because Yang *et al*^[136] showed that in mice injected intraperitoneally with 2.5 \times 10³ *T. gondii* tachyzoites, the serum level of the oxygen free radicals (NO⁺, *OH, O₂⁻) increased along with the days of infection, and superoxide dismutase concentration reached a peak on the 3rd day of the infection. It was documented that oxidative stress with markedly increased serum MDA and decreased glutathione levels is characteristic for chickens orally infected with tissue cysts of the protozoan^[133] (**Tables 6 and 7**), and *T. gondii* seropositive patients (**Table 8**)^[137, 138].

Al-Kennany^[133] demonstrated a lipid peroxidation status in the aortic and heart tissue of *T. gondii*-infected broiler chickens with a significant increase in MDA level and a marked decrease in glutathione concentrations in these tissues. Histopathological examination revealed that aortic lesions had thickening of the wall due to accumulation of lipid vacuoles in *tunica media* extending towards *tunica intima*, associated with proliferation and hypertrophy of VSMCs, hypertrophy of endothelial cells with invasion of tachyzoites in these cells^[133]. Some aortic sections revealed accumulation of tachyzoites in the cytoplasm of VSMCs, well as an accumulation of lipid vacuole concomitant with foamy cells in both *tunica media* and *intima* with fragmentation of elastic *lamellae* and infiltration of lymphocytes^[133] (Table 6).

In asymptomatic *T. gondii*-seropositive blood donors, oxidative stress and immunosuppression have been demonstrated^[139,140]. Foroutan-Rad *et al*^[141] searched in 7 electronic databases information on frequency of toxoplasmosis in blood donors, and in total 43 records which included 20,964 donors from 1980 to 2015, they found that the overall weighted prevalence of exposure was 33% (95% CI, 28-39%). The highest and the lowest seroprevalences of toxoplasmosis were observed in Africa (46%; 95 CI, 14-78%) and in Asia (29%; 95% CI, 23-35%), respectively. Brazil and Ethiopia were identified as countries with high anti-*T. gondii* seroprevalence^[141].

Elsheikha *et al*^[139] found in a group of 260 blood donors that the plasma concentration of MDA was significantly higher and activity of glutathione peroxidase and level of tocopherol $(\alpha, \gamma \text{ and } \sigma)$ fractions (p < 0.001) were lower in *T. gondii*- IgG seropositive than

Table 6 Levels of malondialdehyde (MDA) and glutathione (GSH) in aorta and heart of broiler chickens infected intraperitoneally with 50 tissue cysts of *T. gondii* (acc. to Al-Kennany^[133]; with own modification).

with own modification).							
Groups	MDA (nM per gm wet tissue) Aorta	Heart	GSH (mM per gm wet tissue) Aorta	Heart			
Controls	125 ± 1.05	105 ± 0.33	0.622 ± 0.022 a	0.773 ± 0.026			
Infected	529 8 ± 2 32 a	455.3 ± 0.43 a	0.273 ± 0.01	0.421 + 0.031			

Results are expressed as means \pm SD of 10 broiler chickens per group. ^a Significant differences at $p \le 0.05$.

in seronegative blood donors. The authors suggested that there was a degradation of their antioxidant enzymes caused by oxidative stress induced by increased free radicals attributed to *T. gondii* infection. It must be noted that glutathione induces egress of the parasite from infected cells by activating a *T. gondii*-secreted apyrase (nucleoside triphosphate hydrolase) in the parasite vacuole resulting in a rapid depletion of host cell ATP^[142].

Interestingly, Jafari *et al*^[143] in the case-control study demonstrated that women were more sensitive to the effects of *T. gondii* infection on oxidative stress induction compared to men. Overall, 75 (50%) of 150 participants 20-50 yrs old were seropositive for anti-*T. gondii* IgG antibodies, including 23 (15.33%) men and 52 (34.66%) women; the control group of participants consisted of 75 individuals (32 men and 43 women) with serum anti-*T. gondii* IgG negative samples. Serum catalase activity (converts H_2O_2 to H_2O_1) was significantly increased in men of the case group (p = 0.006), while it was decreased in women of this group, as compared with controls (p = 0.043). Catalase activity in women of the case group was markedly lower than that of men (p = 0.003)^[143]. Effects of *T. gondii* infection on serum glutathione and MDA levels in the case and control groups were presented in **Table 9**.

Vegeto $et\ al^{[1:44]}$ showed that estrogens may block iNOS activity and accumulation in LPS-activated microglia cells, as well as reduce the accumulation of nitrites and nitrates consequent to various inflammatory stimuli, and thus play a protective role against neurodegeneration.

Oxidative stress plays an important role also in the pathogenesis of neurodegenerative diseases. Dincel and Atmaca^[145] demonstrated in the brain tissue of mice infected with ME49 strain *T. gondii* that

Table 8 Serum glutathione, malondialdehyde, and NO concentrations in *T. gondii*-seropositive patients and healthy controls (acc. to Karaman *et al*^[137]; with own modification.

Bioparameter	Group	No of participants	Serum levels (mean ± SD)	P values
Glutathione	Patients	37	3.96 ± 0.10	0.001
Giutatnione	Controls	40	10.37 ± 0.13	
N. 1. 1: 1.1.1. 1	Patients	37	41.32 ± 2.05	0.001
Malondialdehyde	Controls	40	9.18 ± 1.21	
NO	Patients	37	47.47 ± 1.00	0.001
NO	Controls	40	39.18 ± 1.29	

Results statistically significant at p < 0.05. Serum glutathione and NO levels are expressed as μ M, and malondialdehyde concentrations represent nM.

Table 9 Effects of *T. gondii* infection on serum glutathione (GSH; an antioxidant) and malondialdehyde (MDA - a lipid peroxidation marker) levels in anti-toxoplasma IgG-seropositive participants and IgG-seronegative controls (acc. to Jaffari *et al*^[143]; with own modification).

Parameters	Con	trols	0	- seropositive cipants
	Men (n = 43)	Women (n = 32)	Men (n = 23)	Women (<i>n</i> =52)
Weight (kg)	36.744 ±1.185	34.437±1.382	39.565±1.415	36.077±0.779
GSH (nM protein)	0.838±0.059	0.891±0.046	0.757±0.075	0.622±0.053 a
MDA (nM protein)	1.669±0.099	1.902±0.086	1.832±0.107	2.305±0.095 b,c

Results are mean \pm SE. ap = 0.032 and bp = 0.007 vs the controls; cp = 0.028 vs infected men.

Table 7 Plasma lipid profile of broiler chickens infected intraperitoneally with 50 tissue cysts of T. gondii (acc. to Al-Kennany [25] with own modification),

		I		(0.001.1		
Groups	Total cholesterol mg/dL	Triglycerides mg/dL	HDL-C mg/dL	LDL-C mg/dL	VLDL-C mg/dL	Atherogenic index
Controls	117 ± 0.23	105.3 ± 1.28	42.06 ± 2.72 a	53.88 ± 1.25	21.06 ± 0.02	2.78 ± 0.12
Infected	478 ± 0.30 a	192.3 ± 0.48 a	33.36 ± 1.04	406.81 ± 1.04 a	38.49 ± 1.28 a	14.17 ± 0.81 a

Results are expressed as means \pm SD of 10 broiler chickens per group. ^a Significant differences at $p \le 0.05$. HDL-C, high density lipoprotein-cholesterol; LDL-C, low density lipoprotein-cholesterol; VLDL-C, very low density lipoprotein-cholesterol.

the levels of glutathione reductase (p < 0.005) and neuron specific enolase (p < 0.001) expression markedly increased, while Cu/Zn superoxide dismutase activity decreased (p < 0.001) compared to the noninfected animals. Moreover, intense staining for 8-hydroxy-2'-deoxyguanosine (a critical biomarker of oxidative stress) (p < 0.05) was observed both in the nucleus and the cytoplasm of neurons and glial cells that underwent oxidative stress^[145].

Together, there is the vicious circle between oxidative stress and inflammation in atherosclerosis^[104]. Latent chronic *T. gondii* infection play an important role in inducing local/systemic inflammation and oxidative stress in both animals and humans. Since the parasite attacks almost all nucleated cells in mammals and slowly is becoming a global hazard^[30], it seems that it have a pivotal role in development of atherosclerosis.

IMPORTANT ROLE OF T. gondii INFECTION IN THE PATHOGENESIS OF ATHEROSCLE-ROSIS

The parasite has the autonomous capacity to synthesize phospholipids, and also readily scavenges precursors of the lipids from the host cell^[146-148]. *T. gondii* is unable to synthesize sterol and acquires cholesterol from the host LDL receptor (LDLR) pathway^[149]. LDLs are internalized by the LDLR or scavenger receptors and delivered to the late endosomes/lysosomes for hydrolysis^[148]. Deposition of excess cellular cholesterol in the form of CEs is

Table 10 Uptake of cholesterol by mouse embryonic fibroblasts (MEF) ACAT1 $^{-/-}$ cells transfected with TgACAT1 or TgACAT2, and incorporation into cholesteryl esters (acc. to Lige $et~al^{[153]}$; with own modification).

MEF ACAT ^{-/-}	Cholesterol uptake	Cholesterol ester levels
transfected with	(cpm x µg cell protein ⁻¹)	(cpm x µg cell protein-1)
Vector	3991 ± 601	ND
TgACAT1	4003 ± 510	898 ± 65 ^a
TgACAT2	3758 ± 436	1514 ± 110 ^b

The results are mean values \pm SD from three incubations ("a" vs "b" is statistically significant, p < 0.01). MEF ACAT' were transfected with vector alone, or plasmids containing either TgACAT1 or TgACAT2. Cells were then incubated with [3H]cholesterol-LDL. Cholesterol uptake was monitored by scintillation counting. Cholesterol ester detection was performed by thin-layer chromatography analysis. Cpm, counts per minute; ND, not detected.

Table 11 Comparison of free fatty acids (FFAs) uptake by T. gondii and CHO cells (acc. to Quittnat $et\ al^{[156]}$; with own modification).

	Oleate	Palmitate	Stearate	Linoleate	Arachidonate
T. gondii	780 ± 99	1833 ± 105 a	810 ± 69	750 ± 91	767 ± 77
CHO cells	1230 ±142	1156 ± 101	1467 ± 151	1350 ± 98	1058 ± 103

Results are means \pm SD of four separate experiments. Differences between values of the uptake of palmitate vs other FFAs in T. gondii are statistically significant. ap < 0.005. CHO, Chinese hamster ovary. Extracellular parasites or CHO cells were incubated for 1 h at $37^{\circ}\mathrm{C}$ with the indicated 0.3 mM radioactive FFAs. After washing, the total FFAs uptake was determined by measuring the cell-associated radioactivity and expressed in nM per mg cell protein/h.

catalyzed by the resident ACAT, leading to production of lipid bodies^[150,151].

Sakashita *et al*^[152] demonstrated that immature human macrophages expressed only ACAT1, but the fully differentiated macrophages expressed both ACAT1 and ACAT2, and revealed the presence of these two enzymes mRNAs in human atherosclerotic aorta. Immunohistochemical study indicated that in the atherosclerotic aorta all macrophages expressed ACAT1, while approximately 70-80% of the cells also expressed ACAT2. It was suggested that in atherosclerotic plaque, the ability of macrophage foam cell transformation may be enhanced by the dual expression of ACAT1 and ACAT2^[152]. It must be emphasized that the marked expression of these two enzymes present in macrophages may be further augmented by the superimposing *Tg*ACAT1 and *Tg*ACAT2 activities available in the cells infected with *T. gondii* (**Table 10**)^[153].

T. gondii expresses two CE-synthesizing enzymes TgACAT1 and TgACAT2 that localize to the parasite ER, and TgACAT2 shares 56% identity with TgACAT1^[153,154]. Both these enzymes contribute to the CEs formation for storage in lipid bodies, and while TgACAT1 preferentially utilizes palmitoyl-CoA, TgACAT2 produces more CEs because it has a broader fatty acid specificity^[154-156]. The parasite was competent to synthesize CEs using host cell-derived cholesterol and fatty acid, and the production was proportional to the amount of lipid taken up by the protozoan over the time^[154,157]. It was demonstrated in vitro that perfusion of LDL in the culture medium lead to massive uptake of cholesterol by the parasite, suggestive of uncontrolled uptake of this lipid by T. gondii^[153]. It should be emphasized that ACAT and CEs play a crucial role in replication of the pathogen^[157].

It appeared that several FFAs were incorporated into parasite CE but nearly twofold higher incorporation of palmitate into CE as compared with oleate, arachidonate, stearate, and linoleate was observed^[154,156] (**Tables 11 and 12**). The preferential palmitate incorporation specificity by *T. gondii* acyl-CoA: diacylglycerol acyltransferase (*TgDGAT1*) has already been found also for triacylglycerol synthesis in the parasite^[158] (triacylglycerols are quantitatively the most important storage form of energy for eukaryotic cells), and *TgDGAT1* is an integral membrane protein localized to the parasite cortical and perinuclear endoplasmic reticulum^[156]. The protozoan was able to take up both oleate and diacylglycerol and incorporate them into the triacylglycerol fraction, and oleate was incorporated into diacylglycerol (**Table 13**).

Table 13 Lipid acquisition and substrate incorporation into n-acylglycerides of *T. gondii* (acc. to Quittnat *et al*^[156]; with own modification).

	Lipid uptake (cpm per µg cell protein)	Lipid synthesis (cpm per µg cell protein) DAG	TAG
[3H]Oleate	78589 ± 190	10444 ± 1874	32400 ± 3880
[14C]DAG	58430 ± 3209	39650 ± 4705	23664 ± 2099

The values are means ± SD of three separate experiments. DAG, diacylglycerol; TAG, triacylglycerol; cpm, counts per minute. Extracellular *T. gondii* were incubated for 1 h at 37°C with radioactive oleate or DAG before measurement of total lipid associated to parasites and lipid incorporation into DAG or TAG.

Table 12 Fatty acid specificity for triacylglyceride (TAG) synthesis in *T. gondii* (acc. to Quittnat *et al*^[156]; with own modification).

	Relative activities [³H]oleate + oleate	[³H]oleate + palmitate	[³H]oleate + stearate	[³H]oleate + linoleate	[³H]oleate + arachidonate
T. gondii	1.00	0.18	0.69	0.50	0.61
CHO cells	1.00	0.52	0.62	0.55	0.59

The values expressed as relative activities, are means of three separate experiments. CHO, Chinese hamster ovary. Extracellular *T. gondii* and CHO cells were incubated for 1 h at 37 °C with 10 nM of radioactive oleate (~50,000 dpm per nM) previously mixed with 0.3 mM of the indicated unlabeled FFAs to measure its incorporation into TAG. The production of TAG with oleic acid alone was 45,950 cpm per mg cell protein for *T. gondii*.

In atherosclerotic patients as well as during chronic latent T. gondii infection there is an increased production of proinflammatory cytokines that enhances both ACAT1 and ACAT2 activity in the infected host liver hepatocytes, intestine cells, and macrophages^[159-163]. The enhanced activity of these enzymes will thus be overlapping with TgACAT1 and TgACAT2 activities in the intracellular tachyzoites. As a result, there will be an accumulation of a marked surplus of these enzymes activity finally leading to a more efficacious generation of foam cells, and development of several liver and/or intestine abnormalities, such as steatohepatitis or clear cell $colitis^{[164,165]}$. These cells morphologically represent cholesterol ester-laden macrophages/foam cells, because the number and the size of lipid bodies in T. gondii vary between the different parasite developmental stages (tachyzoites, bradyzoites, oocysts), environmental conditions[153,166], infected cells/tissues of the host, and intensity of oxidative stress associated with the infection.

Atherosclerosis was found to be more common in children with fatty liver (30%) than in those without this abnormality (19%)[167,168]. Nonalcoholic fatty liver disease (NAFLD) is the most common of chronic liver disease in Western countries that affects about 20-30% of the general population and is closely related to oxidative stress and insulin resistance[164,169-171]. The prevalence of NAFLD was about 3-10% in lean children, and about 53% in obese pediatric population^[172]. Its histologically proven prevalence in children in the US (as found at autopsy after accidents) ranges from 9.6% in normalweight individuals up to 38% in obese ones^[173]. This clinical entity includes a wide spectrum of liver diseases ranging from steatosis alone, to nonalcoholic steatohepatitis, which may progress to liver fibrosis and cirrhosis[170]. NAFLD is associated with coronary artery disease and subclinical atherosclerosis, emphasizing the role of chronic inflammation in the pathogenesis of atherosclerosis^[171]. Autopsy findings in 817 children who died of external causes showed that fatty liver was present in 15% of the children, mild atherosclerosis in 21%, and moderate to severe atherosclerosis was observed in 2%[172]. These findings emphasize the importance of latent chronic T. gondii infection as a potential but so far neglected cause of atherosclerosis sometimes associated with unexplained liver damage and other clinical entities involving this organ, because this widely disseminated pathogen slowly becomes a global health hazard[30,164].

In summary, it was demonstrated that ACAT and CEs play a crucial role in replication of *T. gondii*. Immature human macrophages expressed only ACAT1, but the fully differentiated macrophages expressed both ACAT1 and ACAT2, and revealed the presence of these two enzymes mRNAs in human atherosclerotic aorta. Moreover, immunohistochemical study indicated that in that aorta, all macrophages expressed ACAT1, while approximately 70-80% of macrophages also expressed ACAT2. In congenital hyperlipidemic mice, ACAT2 was also present in lipid-laden cells of the atherosclerotic plaques. It was suggested that in atherosclerotic plaque, the ability of macrophage foam cell transformation may be augmented by the dual expression of ACAT1 and ACAT2. The marked expression of these two enzymes in macrophages may be further amplified by the TgACAT1 and TgACAT2 of T. gondii present in the infected host macrophages, finally leading to a more efficacious and faster building-up of atherosclerotic plaques.

ACAT1 AND ACAT2 PLAY A CRUCIAL ROLE IN FOAM CELL FORMATION AND DEVELOPMENT OF ATHEROSCLEROTIC PLAQUE.

UNCONTROLLED UPTAKE OF CHOLESTEROL AND SYNTHESIS OF LIPIDS BY *T. gondii* IS DUE TO THE INTRACELLULAR REPLICATION OF THE PARASITE

Atherosclerosis

It is accepted that atherosclerosis involves chronic inflammatory reaction^[175]. The accumulation of foamy transformed macrophages in the aortic intima is a pathological hallmark of the early phase of atherosclerosis^[176,177]. The initial process of foam cell formation is migration of blood monocytes into the vascular wall followed by their differentiation into macrophages, and their conversion into foam cells, which are characterized by intracellular accumulation of cholesteryl esters (CEs)^[178]. In the later stages of atherosclerosis, foam cell undergo apoptosis and secondary necrosis, which causes atherosclerotic plaque rupture^[176]. Complex fibrotic plaques are produced as a result of lysis of foam cells, migration and proliferation of vascular smooth muscle cells (VSMCs), and persistent inflammatory response^[161].

Foamy macrophages bearing numerous lipid droplets are mainly composed of CEs and triglycerides^[6]. These cells secrete various biologically active molecules and promote vascular remodeling with lipid deposition, eventually resulting in pathological vascular events, including myocardial infarction and cerebral ischemia^[176,179,180]. Low density lipoprotein (LDL) is the major cholesterol carrier in the blood and continuous uptake of native and denaturated LDLs by macrophages occurs in the subendothelial layer of the arterial intima^[179].

Important role of ACAT1 in foam cells formation and development of atherosclerotic lesion

Three enzymes are thought to be responsible for the synthesis of plasma CEs, i.e. ACAT1, ACAT2^[176,181,182], and lectin: cholesterol acyltransferase (LCAT), a glycoprotein that is secreted by the liver into the blood^[183]. In animals, ACAT1 and ACAT2 were united with LCAT as known sources of CEs^[184]. In humans, ACAT2 seems to play a much more extensive role than ACAT1 in production of CEs by macrophages because TgACAT2 induced during intracellular T. gondii infection provide additional enzyme activity that further enhances biogenesis of CEs in the host cells^[153,154].

ACAT1 is an essential enzyme in intracellular cholesterol homeostasis and a transmembrane protein localized mainly in the rough endoplasmic reticulum (ER) and perinuclear region that covalently joins excess free cholesterol with fatty acyl-CoA molecule, typically oleoyl-CoA, to form CEs^[185]. ACAT1 and neutral cholesteryl ester hydrolase regulated cholesterol esterification^[2]. Esterification causes cholesterol to be moved out of the cell membrane into the cytoplasm where it is stored as lipid droplets^[184]. The build-up of free cholesterol in membranes can cause membrane deformation, leading to cellular toxicity^[186,187]. The presence of ACAT1 in the cell membrane limits the plasma membrane cholesterol concentration by inducing cytosolic lipid droplets formation. When cholesterol is in overabundance, such as in arterial macrophages during atherogenesis, ACAT1 may promote enough lipid storage to induce conversion of macrophages to lipid-laden foam cells^[184].

ACAT1is a dominant isoenzyme in monocyte-macrophages^[181,188], which plays an essential role in atherosclerotic plaque formation as well as in oxLDL-induced VSMC foam cell build-up^[189,190]. Expression of ACAT1 and ACAT2 significantly increased after differentiation of monocytes to macrophages^[152,191]. In macrophages

and other cell types, ACAT1 overexpression was involved in forming CEs as lipid droplets^[191,192], present in the cell cytoplasm^[177], and finally in foam cell formation^[193,194]. Rong *et al*^[195] found an increase in ACAT1 activity after cholesterol loading also in both human and rat aortic SMCs, but ACAT2 was not found to be expressed by arterial SMCs^[152]. TLR4 has a critical role in triggering inflammation^[189] and participates in VSMC activation^[196]. Unlike the macrophages, VSMCs do not have the proinflammatory properties normally, however, in many atherogenic conditions, inflammatory reaction often was present also in these cells^[197]. It was demonstrated that TLR4-mediated inflammation has been activated by oxLDL in VSMCs, which in turn upregulated ACAT1 expression and finally contributed to VSMC foam cell formation required in atherosclerotic plaque building^[189].

Sakashita *et al*^[198] demonstrated that in normal human tissues ACAT1 was detected in macrophages, smooth muscle cells, antigen-presenting cells, steroid hormone-producing cells, neurons, myenteric ganglia, cardiomyocytes, hepatocytes, mesothelial and other cells, follicles, tubules. The immunohistochemical reactivities were particularly prominent in the macrophages, steroid hormone-producing cells, followed by hepatocytes, and intestinal epithelia. In cultured human macrophages ACAT1 was located mainly in the tubular ER, and on cholesterol loading, about 30 to 40% of the total immunoreactivity appeared in small-sized vesicles enriched in 78 kD glucose-regulated protein^[198].

Miyazaki *et al*^[191] examined human atherosclerotic lesions using immunohistochemical staining and showed that in early lesions of the human aorta, mononuclear cells expressed only limited ACAT1. In contrast, the same ACAT1 antibodies stained much more intensely in fatty streak lesions, particularly in areas that contained macrophages with foamy transformation. It was demonstrated that monocytes/macrophages were the major cellular component of the ACAT1-expressing cells in atherosclerotic lesions^[191]. In animal experiments, cholesterol feeding increased ACAT1 mRNAs by 2- to 3-fold in mouse and rabbit livers and in rabbit aortas^[199,200], respectively, suggesting that the ACAT1 message level may be regulated in a tissue specific manner^[191].

Factors affecting ACAT1 upregulation or downregulation

Cheng et $al^{[201]}$ showed that cholesterol itself served as an activator of ACAT1, besides serving as a substrate. The biosignals that upregulated the expressions of ACAT1 in monocyte-derived macrophages included IFN- γ (a proatherogenic cytokine), dexamethasone, and vitamin $D^{[201-203]}$. On the other hand, a synthetic retinoid Am80 (a specific retinoic acid receptor- α agonist which ameliorated various immunological diseases^[204]), reduced the expression of ACAT1 and HMG-CoA, increased diacylglycerol acyltransferase-1(DGAT-1) expression, inhibited T gondii -induced acquisition and synthesis of cellular cholesterol in macrophage cell line J774.1, suppressed the growth of the parasite, and inhibited IL-6 and atherosclerosis^[205,206].

Insulin was shown to enhance ACAT1 expression in THP-1 macrophages^[207]. Leptin, an adipose tissue-derived hormone, also accelerated CEs accumulation by increasing ACAT1 expression in human monocyte-derived macrophages and cholesterol efflux from these cells^[208]. Hori *et al*^[209] found that TGF-β1, a multifunctional cytokine with important roles in both host defense and immunopathogenesis, increased ACAT1 protein expression and activity by 2- to 3-fold during differentiation of human monocytes into macrophages. It was suggested that ACAT1 expression in monocytes infiltrating from the blood circulation to vascular

walls may be enhanced by pre-existing increased serum TGF-β1 concentrations^[209]. (Nb. TGF-β expression in human retinal pigment epithelial cells enhanced by *T. gondii* may play an important role in the pathogenesis of retinochoroiditis^[210]). Infection with *Chlamydia pneumoniae* (a pathogen isolated from coronary and carotid atheromatous plaques which plays an important role in cardiovascular disease^[211], hides inside apoptotic neutrophils to silently infect and propagate in macrophages^[212]) also induced macrophage-derived foam cell formation by upregulating ACAT1 expression and thus disturbed cholesterol homeostasis in these cells^[213,214]. At present, one cannot exclude also a pathogenic role of *Helicobacter cinaedi* (one of the the most common enterohepatic *Helicobacter* species that causes bacteriemia in humans) infection in promoting atherosclerosis by involving macrophage-driven proinflamatory responses associated with TLR2 and TLR4 signaling, and causing foam cell formation^[215].

Interestingly, Zhao *et al*^[216] showed that hydrogen sulfide (H₂S), a novel endogenous gasotransmitter and a well-known toxic gas with a characteristic smell of rotten eggs present also in drinking waters of wells and thermal spring and mineral waters, inhibited human monocyte-derived macrophage foam cell formation by suppressing ACAT1 production. H₂S shares similar features with NO, and CO, and play important vasodilatatory and antiatherosclerotic roles^[216-218]. These cytoprotective actions may at least partly explain H₂S beneficial effects in slowing down progression of experimental Alzheimer's disease^[219], as well as improving health in major cardiovascular diseases^[220]. It was recently suggested that atherosclerosis and Alzheimer are diseases with a common cause^[221], and it seems that chronic *T. gondii* infection may be at least in part responsible for this causal relationship^[222,223].

Crucial role of ACAT2 in foam cells formation and development of atherosclerotic lesion

ACAT2 is expressed only in intestinal enterocytes, hepatocytes (where ACAT1 is silent[^{224}]), and in macrophages[^{152,225}]. Parini *et al*^{1226} found that ACAT2 protein expression was localized to hepatocytes as the major cholesterol-esterifying enzyme in human liver, whereas ACAT1 was present in Kupfer cells only. Human fetal liver (20 weeks post-conception) had abundance of ACAT2 mRNA[^{227}], and in congenital hyperlipidemic mice ACAT2 was present in lipid-laden cells of the atherosclerotic plaques[^{152}]. It must be noted that the ACAT2 protein expressed in Chinese hamster ovary cell had a half-life > 6 hrs, whereas the ACAT1 protein had a $t_{1/2}$ of only \cong 20-30 min, suggesting greater stability of ACAT2[^{226}].

ACAT2 participates in: (a) the enterohepatic recirculation of cholesterol, as revealed by the reduced intestinal absorption efficiency in ACAT2-deficient mice^[228]; (b) in regulation of cholesterol metabolism within the hepatocyte; and (c) in CE secretion and transport in plasma lipoproteins^[184].

ACAT2 functions in free cholesterol esterification during cholesterol absorption in the enterocyte^[224]. This is a key cholesterol esterification enzyme necessary for the development of hypercholesterolemia^[228]. Most of the absorbed cholesterol is esterified by ACAT2 for incorporation into chylomicron particles, so that 75-80% of newly absorbed cholesterol transported into the body in chylomicrons as CE^[229]. The metabolism of chylomicrons in the circulation leads to the formation of remnant lipoproteins that retain the CE, which is ultimately delivered to the hepatocytes in the liver^[224]. The LCAT derived CE of HDL is removed from plasma primarily by the liver through selective CE uptake^[230] and by whole HDL particle uptake^[231]. Normally, more than 70% of the cholesterol in LDL and VLDL is esterified, and two enzymes ACAT2 and

LCAT (a liver-derived plasma protein) responsible for the synthesis of plasma lipoprotein CEs^[232] are important determinants of both LDL and VLDL cholesterol concentrations and the subsequent development of atherosclerosis^[233,234]. Lee et al^[233] established that while ACAT2 contributed CEs to newly secreted VLDL, LCAT added CE during LDL particle formation in mice^[224,233]. ACAT2 deficiency led to a marked decrease in the percentage of CEs $(37.2 \pm 2.1\% \text{ vs } 3.9 \pm 0.8\%)$ in plasma VLDL, with a concomitant increase in the percentage of triglyceride (33.0 \pm 3.2% vs 66.7 \pm 2.5%)[233]. LDL is the major cholesterol carrier in the blood, and in early stages of atherosclerosis, continuous uptake of naive or denatured LDLs by macrophages occurs in the subendothelial layer of the aortic intima. The build-up of free cholesterol in membranes can cause membrane deformation, leading to cellular toxicity[179,186,187]. Plasma CEs provided by ACAT2 and LCAT have opposite atherosclerotic potential^[232]. Saturated and monounsaturated CEs derived predominantly from ACAT2 were associated with increased atherosclerosis^[235], whereas polyunsaturated fatty acids containing CEs derived from LCAT were associated with decreased atherosclerosis^[236]. LCAT acts primarily on plasma HDL to produce cholesterol linoleate and arachidonate^[237], and for example, in patients with coronary heart disease, the percentage of cholesterol linoleate was found to be diminished^[238,239]. This is in agreement with the finding of Degirolamo et al^[240] that polyunsaturated fat provide atheroprotection, at least in part, because it limited the accumulation of cholesteryl oleate in favor of cholesteryl linoleate in plasma lipoproteins. In addition, several data suggested that the cholesteryl oleate and cholesteryl palmitate synthesized by ACAT2 are particularly atherogenic[233,235,241] presumably because of the limited ability of macrophages to mobilize cholesterol from these $\mbox{\rm CEs}^{\mbox{\tiny [242]}}.$ It was also proposed that cholesteryl oleate should be used as a biomarker for atherosclerosis^[243].

ACAT2 is an ER-localized transmembrane protein that transfers the fatty acyl moiety of acyl-CoA to free cholesterol producing mainly cholesteryl oleate and palmitate, and ACAT2-derived CEs can be packaged directly into nascent apoB-containing lipoproteins or stored as neutral droplets in the cytosol^[181,182,226,244,245]. Liang et al^[246] demonstrated that overexpression of either ACAT1 or ACAT2 in rat hepatoma cells increased the secretion and cellular accumulation of CE to similar extent, but ACAT2 expression caused a greater increase in apoB secretion than ACAT1. It is believed that the primary role of liver ACAT2 expression, relevant to atherogenesis, was to enrich apoB-containing lipoproteins with cholesteryl oleate and palmitate, as opposed to increasing the number of secreted apoB-containing particles^[234]. The percentage of lipoprotein cholesteryl oleate and the rate of hepatic cholesteryl oleate secretion were found to be positively related to the extent of atherosclerosis in monkeys^[235,247]. Temel et $al^{[234]}$ demonstrated that the relative level of ACAT2 expression in cells affects the CEs content and, hence, the potential atherogenicity, of nascent apoB-containing lipoproteins (Table 14).

Cases *et al*^[155] demonstrated in H5 insect cells (these cells have very low levels of cellular cholesterol and lack meaningful cholesterol esterification activity^[192]) that of the fatty acids tested, ACAT1 showed a slight preference for oleoyl, while that for ACAT2 was palmitoyl \geqslant linoleoyl > arachidonyl (**Table 15**). They also found that for ACAT1, palmitoyl, linoleoyl, and to a lesser extent, arachidonyl, competed with oleoyl for incorporation into CEs, while for ACAT2, both linoleoyl and palmitoyl competed with oleoyl for incorporation into these structures (**Table 16**). Moreover, it was established that ACAT1 had 58% higher activity with oleoyl-CoA compared with palmitoyl-CoA, while ACAT2 showed only small

Table 14 Effect of ACAT2 on the radiolabelled lipid composition of apoB-34-containing lipoproteins (acc. to Temel *et al*^[234]; with own modification).

ACAT2 Plasmid (ug)	Percentage total [³H] lipid						
ACAT2 Plasmid (μg)	PL	TG	CE	CE/TG	PL/CE+TG		
0	19.1	77.7	3.4	0.042	0.24		
1	19.2	68.7	14.5	0.211	0.2		
2.5	19.0	63.5	17.4	0.274	0.24		
5	20.3	57.8	22.0	0.38	0.25		

Values shown are means of duplicate samples. ApoB, apolipoprotein B, CE, cholesteryl ester; PL, phospholipids; TG, triglyceride.

Table 15 Fatty acyl-CoA specificity for cholesterol esterification by ACAT1 and ACAT2 expressed in H5 cells. Esterification of [14 C] cholesterol using different fatty acyl-CoAs as substrates (acc. to Cases *et al*^[155]; with own modification).

Experiment	Relative Oleoyl-CoA	activities (units) Palmitoyl-CoA	Linoleoyl- CoA	Arachidonyl- CoA
ACAT1				
Experiment 1	1.0	0.9	0.7	0.4
Experiment 2	1.0	0.6	0.7	0.8
ACAT2				
Experiment 1	1.0	1.4	1.4	0.6
Experiment 2	1.0	1.7	1.4	0.7

Cell membranes were prepared and assayed in triplicate. The cholesterol: phospholipid liposome molar ratio was 0.2. For ACAT1, the values with oleoyl-CoA were 566 and 499 pM of cholesterol esters formed/mg protein/min for experiments 1 and 2, respectively, and for ACAT2, they were 594 and 529 pM of cholesterol esters formed per mg of protein/min.

Table 16 Fatty acyl-CoA specificity for cholesterol esterification by ACAT1 and ACAT2 expressed in H5 cells. Competition of fatty acyl-CoAs with $[^{14}C]$ oleoyl-CoA for cholesterol esterification (acc. to Cases et $al^{[155]}$; with own modification).

Experiment	Relative Oleoyl-CoA	activities (units) Palmitoyl-CoA	Linoleoyl- CoA	Arachidonyl- CoA			
ACAT1							
Experiment 1	1.0	0.7	0.8	0.9			
Experiment 2	1.0	0.8	0.8	0.8			
ACAT2	ACAT2						
Experiment 1	1.0	0.8	0.7	1.3			
Experiment 2	1.0	0.8	0.8	1.3			

Cell membranes were prepared and assayed in triplicate. The cholesterol: phospholipid liposome molar ratio was 0.7. For ACAT1, the values with oleoyl-CoA were 749 and 644 pM of cholesterol esters formed/mg protein/min for experiments 1 and 2, respectively, and for ACAT2, they were 1661 and 1590 pM of cholesterol esters formed per mg of protein/min.

Table 17 Fatty acyl-CoA specificity for cholesterol esterification by ACAT1 and ACAT2 expressed in H5 cells. Comparison of cholesterol esterification using [14 C]oleoyl-CoA or [14 C]palmitoyl-CoA (acc. to Cases *et al*] 155 ; with own modification).

Experiment	[14C]Oleoyl-CoA pM cholesterol ester/mg protein/min	[¹⁴ C]Palmitoyl- CoA
ACAT1	529	334
ACAT2	1823	1573

Cell membranes were prepared and assayed in triplicate. Values were \pm < 5% in all cases. The cholesterol:phospholipids liposome molar ratio was 0.7

difference (**Table 17**)^[155]. All these data strongly suggested that ACAT2 play a much more extensive role than ACAT1 in generation of CEs by macrophages^[155]. Miller *et al*^[248] proposed that CEs

associated with ACAT2 may have a predictive value in diagnosis of coronary artery disease in patients with acute chest pain.

Further studies of Seo *et al*^[190] also demonstrated that incubation of oleic acid, arachidonic acid, and eicosapentaenoic acid induced ACAT1 mRNA levels 1.5-2-fold in HepG2 cells, with no effect on ACAT2 mRNA. Both HepG2 and THP1 cells showed the greatest CE production with oleic acid. ACAT1 expressing cells indicated a strong preference for oleic acid while ACAT2 expressing cells utilized unsaturated free fatty acids (FFAs). THP1 and ACAT1 expressing cells utilized oleoyl CoA preferentially. In contrast, HepG2 and ACAT2 microsomes utilized linolenoyl CoA as well. The authors believed that FFAs increased ACAT1 mRNA levels in a cell specific manner, and that the ACAT reactions exhibited differential FFA utilization^[190].

Factors affecting ACAT2 downregulation

Estrogen have beneficial effects on plasma lipids and lipoproteins, and reduced atherosclerosis progression [249,250]. Kavanagh $et~al^{[250]}$ found that estrogen decreased atherosclerosis primarily by reducing hepatic ACAT2 activity in ovariectomized monkeys. Estrogen regulation of enzymatic activity was established at the protein level as both ACAT1 and ACAT2 protein (but not mRNA levels), were lower (p = 0.02 and < 0.0001, respectively)^[250]. Interestingly, hepatic microsomal ACAT2 activity in normolipidemic, non-obese Chinese females was significantly lower than observed in males^[251]. Moreover, the activity of ACAT2 correlated negatively with plasma levels of HDL-C (r = -0.57, p < 0.05) and with Apo AI (r = -0.49, p < 0.05). The observed sex-related difference may thus contribute to female protection from complications of coronary heart disease^[251]. Saba et al^[252] showed that black ginseng extracts (Ginseng Radix nigra, a well-characterized medicinal herb with antioxidant activity and increased radical scavenging properties) significantly downregulated ACAT2 and HMG-CoA mRNA expressions in murine hepatocytes. Chronic fluticasone propionate (a glucocorticoid) exposure also decreased ACAT2 gene expression in human monocytes and macrophages^[253]). Orange juice or hesperidin consumption resulted in a decrease in the expression of genes encoding ACAT2 together with an increase in the expression of genes regulating reverse cholesterol transport[254]. ACAT2 was also downregulated by naringenin, a flavonoid from grapefruit, which could reduce foam cell formation^[254,255]. Interestingly, curcumin (no effect on ACAT1 or ACAT2; inhibition of oxLDL uptake and promotion of cholesterol efflux from macrophages)[256,257] and erythropoietin (increased cholesterol efflux from macrophages)[258], were reported to suppress the formation of macrophage foam cells. Finally, it should be noted that delta 9-tetrahydrocannabinol (THC), the main constituent of marijuana (Cannabis sativa), administered by a noninhalation route may exert dose-dependent (1 mg/kg/day; low 0.6 ng/mL plasma levels with no psychotropic effects) atheroprotective effects via its immunomodulatory and antiinflammatory actions[259], as well as inhibition of ACAT1 and ACAT2 activities^[260].

In summary, ACAT2 enzyme has a longer half-life than ACAT1. For ACAT2, both linoleoyl and palmitoyl competed with oleoyl for incorporation into CEs, while for ACAT1, palmitoyl, linoleoyl, competed with oleoyl for this biochemical reaction. ACAT1 had a higher activity with oleoyl-CoA compared with palmitoyl-CoA, while ACAT2 showed only small difference. All these data strongly suggest that ACAT2 play a much more extensive role than ACAT1 in generation of CEs by macrophages, and thus in foam cells biogenesis. Clinically, it was proposed that CEs associated with ACAT2 may predict coronary artery disease in patients with

acute coronary syndrome, especially that gender-related differences in hepatic microsomal ACAT2 activity have been documented in normolipidemic, non-obese Chinese females.

IMPORTANT ROLE OF TOLL-LIKE RECEPTORS (TLRs), ESPECIALLY TLR2 AND TLR4, IN BOTH T. gondii INFECTION AND DEVELOPMENT OF ATHEROSCLEROSIS

Atherosclerosis

The first steps of atherosclerosis are inflammatory in nature^[40,261]. A family of TLRs has been defined as a key component of pathogenassociated molecular pattern recognition machinery[262,263]. TLRs are proinflammatory sensors of pathogens, and potential links between infectious disease, inflammation and atherosclerosis, originally identified as receptors that activate host defenses in response to microbial-derived ligands such as Gram-negative bacterial lipopolysaccharides (LPS)[264,265], and T. gondii[266,267]. There are growing evidence showing the contribution of the TLR-signaling pathway to initiation and progression of atherosclerosis^[268,269]. Xi et al^[270] performed a systemic analysis of the molecular pathomechanism underlying atherosclerosis and found that 1312 genes from 45,304 publications, were associated with atherosclerosis. 35 gene ontology terms were significantly enriched, and the most highly overrepresented pathway went to the TLR signaling pathway, known to play an important role in atherosclerosis in both inflammatory and immune response[270].

Two prominent risk factors, infectious disease and hyperlipidemia point to innate immune mechanisms as potential contributors to proatherogenic inflammation^[271]. Atherosclerotic lesions form only in arteries at sites of disturbed blood flow^[272]. Although all endothelial cells are exposed to comparable levels of circulating plasma cholesterol, only endothelial cells overlaying lesions display an inflamed phenotype^[75]. This occurs even in the absence of any additional exacerbating disease factors because blood flow controls the expression of TLRs which are initiators of cellular activation and inflammation. TLR2 and TLR4 expression exerted an overall proatherogenic effect in hyperlipidemic mice. TLR activation of the endothelium promoted lipid and leukocyte accumulation within lesions^[75].

TLRs ligands are expressed in various cell types within the atherosclerotic plaque^[273]. TLR1, TLR2, and TLR4 were upregulated in human atheroma with active NF-κB colocalizing with TLR2 and TLR4 in the plaque^[263]. Tobias and Curtis^[271] observed that proatherogenic TRL2 responses to so far unknown endogenous or unknown endemic exogenous agonists were mediated by non bonemarrow-derived cells, including endothelial cells. TLR ligands potentially associated with development of atherosclerosis are presented in **Table 18**.

TLR-2 ligands (lipoproteins, such as peptidoglycan and lipoteichoic acids from Gram-positive bacteria) and TLR-4 ligands (LPS from Gram-negative bacteria) play an important role in the innate and inflammatory signaling to activate the NF-κB pathway and release of proinflammatory cytokines^[276-283]. Activation of NF-κB is essential for the regulation of a variety of genes involved in the inflammatory and proliferative responses of cells critical to atherogenesis^[284,285]. Edfeldt *et al*^[263] found that the expression of TLR1, TLR2 (peptidoglycan as a ligand), and TLR4 (LPS as a ligand) was markedly enhanced in human atherosclerotic plaques and associated with inflammatory activation of endothelial cells and macrophages. TLR expression

by macrophages permitted local differentiation of these cells into antigen-presenting cells because TLR2 ligation stimulated this process^[286]. Therefore, TLRs may provide an important link between local innate and adaptive immunity and enhance the cellular immune response in the plaque to local antigens such as heat shock proteins (HSPs) and oxidized lipoproteins^[263].

In the advanced atherosclerotic lesions in humans, TLR1, TLR2, TLR4 were detected in macrophages, endothelial cells, and vascular smooth muscle cells (VSMCs), while TLR2 and TLR4 were expressed in atherosclerotic lesion^[263,264,280]. Human VSMCs were reported to express functional TLR2 and TLR4 signal complexes linked to chemokine and proinflammatory cytokine release^[287]. The expressions of TLR2 and TLR4 in VSMCs were augmented in the atherosclerotic lesions^[288]. Local overexpression of TLR-like receptors at the vessel wall induced atherosclerotic lesion formation. Shinohara et al^[264] demonstrated that the expression of both TLR2 and TLR4 at the vessel wall, particularly in VMSCs, synergistically accelerated atherosclerotic process at its early stage, especially under hypercholesterolemia. They also showed that overexpression of TLR2 and TLR4 in VMSCs induced the synergistic activation of NFκB at the vessel wall, which resulted in augmented expression of key molecules of atherosclerosis such as intracellular adhesion molecule 1, vascular cell adhesion molecule 1, and monocyte chemoattractant protein-1[264].

Xu et al^[280] showed that the proinflammatory signaling receptor TLR4 colocalized with lipid-rich and macrophage-infiltrated murine atherosclerotic lesions and at the shoulder region of human coronary artery plaques. Furthermore, basal TLR4 mRNA expression by cultured human monocyte-derived macrophages was upregulated by oxLDL (but not native LDL) in vitro, raising the possiblity that enhanced TLR4 expression may play a role in the inflammatory responses linking lipids to chronic infection, inflammation, and atherosclerosis^[40,289,290]. In fact, TLR4 was expressed by macrophages in murine and human lipid-rich atherosclerotic plaques and upregulated by oxLDL^[280].

HSP60 was found to be implicated in the pathogenesis of atherosclerosis^[291], and the intensity of its expression correlated positively with the atherosclerotic severity^[292,293]. Hypoxia induced HSP expression in human coronary artery bypass grafts^[36]. Berberian et al[294] reported also the elevated expression of HSP70 in human atherosclerotic plaques, and it was mainly concentrated in the central portions of more thickened atheromas around sites of necrosis and lipid accumulation[295]. OxLDL triggered in vitro the expression of HSP60 in monocytes/macrophages^[296], and HSP70 in human endothelial cells^[297] and SMCs^[298]. It appeared that oxLDL induced HSP70 expression mainly in growing endothelial cells, indicating that those cells, present in lesion-prone sites, were more sensitive to oxLDL than were quiescent cells^[299]. HSP70 played a major role in cytokine generation in human oxLDL treated macrophages^[300], and it was demonstrated that plasma HSP70 levels predicted risk of acute coronary syndrome[301], the development of atherosclerosis in patients with established hypertension[302], and had atheroprotective properties[303].

Gonzalez-Ramos *et al*^[304] found that HSP70 increased extracellular matrix (ECM) production by human VSMCs through transforming growth factor-β1 (TGF-β1) up-regulation. TGF-β1 is a vascular profibrotic cytokine that is regulated in part by activator protein 1 (AP-1)-dependent transcriptional mechanisms. HSP70 can interact with the VSMCs, the major producer of ECM proteins, through the TLR4. The authors demonstrated that extracellular HSP70 binds to human aorta smooth muscle cell TLR4, which upregulated the

Table 18 TLR ligands implicated in atherosclerosis (acc. to Hodgkinson & Ye^[273], with own modification).

Te / William Milliam Carrenty.					
TLR	Ligand				
TLR2	Oxidized LDL, hyaluronan, HSP60, peptidoglycan, Apo CIII, HMGB1				
TLR3/9	CMV, viral dsDNA				
TLR4	LPS, modified LDL, fibrinogen, hyaluronan, heparan sulfate, fibronectin EDA, HSP60, tenascin C, β-defensin 2, HMGB1				

Apo CIII, apolipoprotein CIII (Apo CIII induces expression of VCAM-1 in vascular endothelial cells and increases adhesion of monocytic cells); CMV, cytomegalovirus; dsDNA, double stranded DNA; fibronectin EDA, fibronectin extra domain A; HMGB1, high-mobility group box protein 1 (a proinflammatory molecule); LPS, lipopolysacharide. Stimulation with heparan sulfate induced dendritic cells (DCs) to secrete proinflammatory cytokines, such as TNF-α and IL-6, an effect that was TLR4 dependent [275,274]. DCs phenotypically and functionally matured in response to heparan sulfate, an effect that was abrogated when TLR4 was inhibited [275].

AP-1-dependent transcriptional activity of the TGF- $\beta1$ promoter. The TGF- $\beta1$ upregulation increased the expression of the ECM proteins type I collagen and fibronectin. These findings may help in elucidating the pathomechanisms by which HSP70 contributed in the inflammation and fibrosis observed in atherosclerosis^[304].

T. gondii infection

TLRs expressed differentially among immune cells^[305-307] recognize microbial components distinct from host self molecules. Glycosylphosphatidylinositol (GPI)-anchored proteins dominate the surface of the *T. gondii* tachyzoite^[308,309], and GPI synthesis is an essential process for viability of the parasite^[310,311]. GPI from tachyzoites induced TNF-α generation in macrophages via the activation of the transcription factor NF-κB^[312]. Debierre-Grockiego *et al*^[307,311,312] suggested that both TLR2 and TLR4 receptors may contribute in collaboration with other MyD88-dependent receptors (*e.g.* TLR9) in the host defense against *T. gondii* infection through their activations by the GPIs. Interestingly, the native core glycans and the diacylglycerols isolated from purified *T. gondii* GPIs also were able to activate both TLR2 and TLR4, and simultaneous deficiency of TLR2 and TLR4 was required to abrogate the *in vitro* TNF-α production by macrophages exposed to *T. gondii* GPIs^[307].

Denkers *et al*^[313, 314] found that macrophages, which along with DCs are targets of *in vivo* infection, produced cytokines such as TNF- α and IL-12 when activated through TLR4 on the host cell surface^[315,316]. If these cells were first exposed to tachyzoites, they lost their ability to respond to IFN- γ , and TNF- α production was suppressed. Because other biomediators such as antiinflammatory IL-10 were unaffected, carrying the parasite did not make such cells generally dysfunctional. Moreover, only cells containing intracellular pathogens became unresponsive and there was no bystander effect on uninfected cells. Instead, *T. gondii* directly subverted these signaling pathways within host cells probably as a defense mechanism to avoid or delay induction of antimicrobial activity and/or T-cell-mediated immunity during host infection^[316,317].

T. gondii-derived HSP70 stimulated maturation of murine marrow-derived DCs through TLR4^[60]. Under stress conditions, *T. gondii* RH strain tachyzoites differentiated into bradyzoites and induced *T. gondii* HSP70 (*Tg*HSP70) protein expression, a highly immunogenic protein^[318-321]. Also, reactivation *Tg*HSP70 *in vivo* induced expression of *Tg*HSP70 during bradyzoite to tachyzoite interconversion^[322]. Recombinant (*rTg*HSP70) and natural *Tg*HSP70 stimulated the NO release by peritoneal macrophages via TRL2, MyD888 and IRAK4, but under restimulation, signaling of *rTg*HSP70-induced tolerance

was mediated by TRL4[323].

HSP70 has been shown to be a major immunomodulant antigen in parasitic and bacterial infections, and the preferred target of humoral and cell-mediated immune responses to infection[320,323-326]. Barenco et al[321] found that in mice the TgHSP70 release into the bloodstream was dependent on the death of T. gondii tachyzoites mediated by the host immune response, whereas the increased TgHSP70 expression in the brain depended on the multiplication rate of the parasite. Aosai et al[327] showed that TgHSP70 induced prominent responses in murine B cells derived not only from T. gondii-infected but also from uninfected mice. B220+ spleen cells showed marked proliferation after stimulation with TgHSP70 but neither CD4⁺ nor CD8⁺ population responded. These data indicated that TgHSP70 induced proliferation of B cells but not T cells, and TLR4 was found to be required for TgHSP70-induced B cell activation^[189]. It should be emphasized that TLR4-mediated inflammation promoted foam cell formation of VSMC by upregulating ACAT1 expression[189].

TgHSP70 was shown to deteriorate the host defense by down-regulating NO release of peritoneal macrophages in the *T. gondii*-infected host^[328]. TgHSP70 induced anti-mHSP70 autoantibody formation by B-1 cells in *T. gondii*-infected mice^[329]. It was reported that HSP70 induced DC maturation after binding to immature DC and suggested the abundant receptor expression for HSP70 on immature DC^[317,330]. It is well known that TRL4 is a critical receptor and signal transducer for LPS. LPS downregulated surface expression of the TRL4-MD2 complex on peritoneal macrophages^[331,332]. It was demonstrated that both TgHSP70- and LPS-induced proliferative responses of spleen cells required TRL4 molecules as a receptor^[327].

The mechanisms responsible for the atherogenic modification of LDL may include: oxidation mediated by NOS, myeloperoxidase, and/or 15-lipoxygenase^[333,334]. NO is a potent oxidant produced by both endothelial cells and macrophages that appear to exert both protective and atherogenic effects, depending on the source of production. NO produced by endothelial NOS has vasodilator function and is potentially atheroprotective. In contrast, NO produced via the much higher capacity iNOS in macrophages, serving antimicrobial functions based on its potent oxidative properties, is potentially atherogenic^[334].

NO and its metabolites control a number of host defense functions regulated by activated macrophages, including antimicrobial and tumoricidal activities, implicated in the pathogenesis of tissue damage associated with acute and chronic inflammation [335-337]. It was demonstrated that NO lead to the induction of HSP70 protein and mRNA in cultured SMCs [338] and other cells [339,340]. Mun *et al* [323] found that TgHSP70-induced NO release was dependent on the TLR2/MyD88 signal transduction pathway, and that prior exposure to TgHSP70 induced a tolerance of NO production for subsequent stimulation with TgHSP70 via TRL4 (**Table 19**). TgHSP70 was shown to cause deterioration of the host defense by downregulating NO release by peritoneal macrophages in the T. gondii-infected host [323,329,341].

Together, TLR-like receptors play a critical role in the innate immune responses to microbial-derived ligands, including those of *T. gondii*. TLR2 and TLR4 ligands are important for the inflammatory signaling to activate the NF-κB pathway and release proinflammatory cytokines. In the advanced atherosclerotic lesions in humans, TLR1, TLR2, TLR4 were detected in macrophages, endothelial cells, and VSMCs, while TLR2 and TLR4 were expressed in atherosclerotic lesions. TLR4-mediated inflammation promoted foam cell formation of VSMC by upregulating ACAT1 expression. Activation of TLR2 and TLR4 by GPIs derived from *T. gondii* associated with

the increased production of proinflammatory cytokines, oxLDL, HSP60, HSP70, and NO and its metabolites are all implicated in the pathogenesis of atherosclerosis.

PROINFLAMMATORY CYTOKINES IN-CREASE FOAM CELL FORMATION. LATENT CHRONIC T. gondii INFECTION PERSIS-TENTLY GENERATES PROINFLAMMATORY CYTOKINES AND THUS PLAY A CRUCIAL ROLE IN FOAM CELL BIOGENESIS

Atherosclerosis

Inflammation is a triggering and an aggravating factor in atherogenesis^[40,161]. Elevated plasma levels of proinflammatory cytokines have been demonstrated in patients with established atherosclerotic disease^[162,163]. It is therefore believed that cytokines are key players in all stages of disease and have a profound influence on the pathogenesis of the disease^[342].

TNF- α , a proinflammatory cytokine, participates in every step of the inflammation process^[159,340-343]. High concentrations of TNF- α were found in atherosclerotic lesions, and it was reported that this cytokine regulated multiple genes involved in various stages of atherosclerosis, exhibiting mostly proatherosclerotic properties^[159]. In animal models of atherosclerosis, inhibition of TNF- α reduced the progression of this process^[347]. In healthy males, plasma TNF- α concentrations correlated with early carotid artery atherosclerosis^[348]. TNF- α is secreted by activated macrophages, lymphocytes, endothelial cells, and SMCs^[349]. Lei *et al*^[159] showed that in cultured differentiating human monocytes, TNF- α even at low concentrations (2.5-5 ng/mL), increased the expression of the ACAT1 gene, enhanced the CEs accumulation, and promoted foam cells formation.

Cytokine regulation of ACAT1 gene expression in monocytes/macrophages play important roles in the initiation and progression of atherosclerosis^[159]. The expression of the human ACAT1 gene was also synergistically upregulated by IL-1 β ^[160,161], IFN- γ ^[203,347], and TGF- β ^[210]. In contrast, the human ACAT1 gene expression was downregulated by adiponectin, an adipocytokine that exerted many antiatherosclerotic effects in cell culture studies^[350], for example by increasing cholesterol efflux from macrophages^[351].

Lipid loading of primary human macrophages resulted in altered cytokine secretion from cells and effects were similar regardless of neutral lipid composition of cells^[352]. It was demonstrated that formation of triglyceride (TG)-loaded human monocyte-derived macrophages caused a 4-fold increase in basal IL-1β secretion, and

Table 19 Tolerance of NO synthesis by secondary challenge with TgHSP70 through TLR4 (acc. to Mun *et al*^[323]; with own modification).

Mouse strain	Stimulation	NO synthesis ^a	Tolerance induction ^b
Wild type	TgHSP70	+	+
whatype	LPS	+	+
TLR4 deficient	TgHSP70	+	-
1 LK4 delicient	LPS	-	ND
TLR2 deficient	TgHSP70	-	ND
1 LKZ delicient	LPS	+	+
MyD88 deficient	TgHSP70	-	ND
wiy Doo dencient	LPS	-	ND

a+, NO synthesis positive; -, NO synthesis negative; b+, occurrence of tolerance induction; -, no occurrence of tolerance induction. ND, not detected, NO synthesis did not occur by primary stimulation with TgHSP70 and LPS. LPS, lipopolysaccharide. TLR2 regulated TgHSP70-induced TNF- α mRNA expression and tolerance.

in contrast, secretion of TNF-α and IL-6 decreased significantly following both TG and cholesterol loading[352]. Formation of TG loaded foam macrophages by incubation with VLDL also resulted in a marked increase in IL-1β expression in these cells^[353]. Ruan et al[161] showed that IL-1β (5-20 ng/mL) enhanced transformation of VSMCs into foam cells by increasing uptake of unmodified LDL via LDL receptors and by enhancing cholesterol esterification. It was found that IL-1ß increased both ACAT1 protein activity and mRNA expression. Exposure to IL-1β also caused overexpression of the sterol regulatory element binding protein (SREBP), cleavageactivating protein, and enhanced its translocation from the endoplasmic reticulum to the Golgi, where it is known to cleave SREBP, thereby enhancing LDL receptor gene expression. Thus, IL-1β disrupted cholesterol-mediated LDL receptor feedback regulation, permitting intracellular accumulation of unmodified LDL and causing foam cell formation. This suggested that proinflammatory cytokines may contribute to intracellular accumulation of native LDL without prior modification of the lipoprotein by oxidation or glycosylation under inflammatory stress[161].

IL-33 exerted a controling protective role in atherosclerosis *in vivo*^[354]. This IL-1 cytokine family member induced the production of the T_H2 cytokines IL-4 and IL-13, and enhanced serum immunoglobulin synthesis^[355]. IL-33 was a promoter of T_H2-dependent inflammatory diseases^[356,357], and activated a number of cell types, including T_H2 cells, mast cells and basophils, and promoted systemic sclerosis^[358]. McLaren *et al*^[354] found that IL-33 blocked macrophage foam cell formation *in vitro* by directly decreasing acetylated LDL and oxLDL uptake, storage of cholesteryl esters and triglyceride, and reducing intracellular total and esterified cholesterol content, and enhancing cholesterol efflux/transport, and by inducing a phenotypical T_H1-to-T_H2 switch. IL-33 also decreased the expression of key genes implicated in cholesterol esterification and triglyceride storage, including ACAT1 and adipocyte differentiation-related protein^[354].

Feingold *et al*^[359] found that low doses of LPS stimulated hepatic TG production, while high doses inhibited the clearance of TG-rich lipoproteins in rats. LPS and several cytokines, including TNF- α , TNF- β , IL-1, IL-6, IFN- α , platelet activating factor, rapidly induced *de novo* fatty acid (FA) synthesis and hepatic TG synthesis in rodents (**Table 20**)^[360-362], at least in part by increasing hepatic intracellular concentrations of citrate, an activator of acetyl CoA carboxylase ^[363]. IL-4, an antiinflammatory cytokine, inhibited the stimulatory effects of TNF, IL-1, and IL-6 on hepatic FA synthesis by blocking the increase in hepatic citrate levels^[364]. LPS stimulated hepatic cholesterol synthesis by increasing the transcription rate, mRNA expression, protein mass, and activity of HMG-CoA reductase^[365,366]. In rats, LPS significantly inhibited the clearance of LDL from the circulation^[367], and decreased the expression of hepatic LDL receptor protein (**Table 21**).

T. gondii infection

Acute toxoplasmosis sometimes leads to lethal overproduction of proinflammatory $T_{\rm H}1$ cytokines^[33]. During acute and chronic infection in mice there were differences in virulence of *T. gondii* and different strains of the parasite induced different cytokine responses^[368,369]. For example, the serum levels of IL-12, IL-6, IL-10, IFN-γ, and TNF-α were significantly higher in the type II strain ME49-infected mice than in the FORT-infected mice, suggesting that not all type II strains (most frequent in humans) cause toxoplasma encephalitis in these animals^[368]. Chronic infection with *T. gondii* induced high levels of the cytokines IL-12 and TNF-α, which are

typically associated with macrophage function[316], and DCs[370,371] as well as neutrophils[372] also have emerged as essential producers of IL-12. A pyrogenic factor TNF-α is liable to induce the secretion of acute inflammatory phase proteins via the production of IL-6[373,374]. In toxoplasmosis, TNF-α appeared to be essential for macrophage activation and inhibition of parasite replication, and this action can only be exerted in synergy with IFN-γ^[373]. Macrophages are known as important effectors of the innate immune system, and they posses the potential of directing acquired immunity toward a T_H1-biased type of immune response^[375]. In addition to producing proinflammatory cytokines such as IL-12, TNF-α, IL-1β[32,376-379] and antiinflammatory cytokines (eg, IL-10, TGF-β)[32], macrophages are a source of the microbicidal effector molecule NO and are capable of phagocytosing pathogens and degrading them within the phagolysosome. It should however be noted that the parasite also displays potent downregulatory effects on IL-12, TNF-α, and NO production by infected macrophages, being a result of tachyzoite-induced blockade of NF-κB nuclear translocation^[316,380]. In addition, *T. gondii* readily uses macrophages as host cells and reside in these cells within a specialized parasitophorous vacuole[381,382], which resists acidification and lysosomal fusion[382].

IL-6 and IL-27 are closely related cytokines that play critical but distinct roles during infection with *T. gondii*, i.e. IL-6 is required for the development of protective immunity to this pathogen, whereas IL-27 is required to limit infection-induced pathology^[383]. IL-6 is known to mediate both pro- and antiinflammatory effects having two distinct ways to induce cell-signaling: either through the membrane bound receptor (antiinflammatory) or through trans-signaling (proinflammatory)^[384]. This cytokine is the main stimulator of the production of most acute phase proteins^[385,386], and is important to the transition between acute and chronic inflammation^[387]. Moreover, IL-6 played an important role in regulation of the expression of

Table 20 Effects of LPS, lipoteichoic acid (LTA), and cytokines on triglyceride (TG) metabolism in intact rodents (acc. to Khovidhunkit et $al^{[360]}$; with own modification).

Parameter	LPS	LTA	TNF	IL-1	IL-6	IFN-α	IFN-γ
Serum TG	1	1	1	1	1	\leftrightarrow	\leftrightarrow
Hepatic FA synthesis	1	1	1	1	1	1	\leftrightarrow
FA oxidation	\	ND	\	↓	ND	ND	ND
TG secretion	1	1	1	1	1	ND	ND
TG clearance	\leftrightarrow^a , \downarrow^b	\leftrightarrow	\leftrightarrow	\leftrightarrow	\leftrightarrow	ND	ND
LPL activity	↓	\leftrightarrow	↓, ↔ ^c	\	↓	↓	↓
Lipolysis	1	1	1	\leftrightarrow	1	1	1
Serum ketone body	↓	ND	↓	↓	\leftrightarrow	$\uparrow^a, \leftrightarrow^b$	1

FA, fatty acids; LPL, lipoprotein lipase; LPS, lipopolysacchride; ND, not determined. ^aLow doses. ^bHigh doses. ^cSome but not most tissues.

Table 21 Effects of LPS, lipoteichoic acid (LTA), and cytokines on cholesterol metabolism in intact rodents (acc. to Khovidhunkit $et\ al^{[360]}$; with own modification).

The other mounication,							
Parameter	LPS	LTA	TNF	IL-1	IL-6	IFN-α	IFN-γ
Serum cholesterol	↑,↓ ^a	1	↑,↓ ^a	↑,⇔ ^a	1	\leftrightarrow	\leftrightarrow
Hepatic cholesterol synthesis	1	ND	1	↑	ND	\leftrightarrow	1
HMG-CoA reductase activity	1	ND	1	↑	ND	ND	\leftrightarrow
LDL receptor protein	\downarrow , \leftrightarrow	ND	\leftrightarrow^{b}	\leftrightarrow^{b}	ND	ND	ND
Bile acid synthesis	1	ND	1	\	ND	ND	ND

LDL, low density lipoproteins; HMG-CoA, 3-hydroxy-3-methylglutamyl coenzyme A; LPS, lipopolysacchride; ND, not determined. ^aPrimates. ^bHamsters.

cytokine and chemokine stimulation of antibody production by B cells, regulation of macrophage and DC differentiation, and the response of regulatory T cells to microbial infection [388-390]. This cytokine also protected human macrophages from cellular cholesterol accumulation [391]. Most importantly however, because IL-6 was found to promote the intracellular multiplication of T. gondii in mice [392], and in peritoneal macrophages (in a dose-dependent manner) [393], it may play an important role in triggering and development of atherosclerosis.

Lipoprotein-associated phospholipase A₂ (Lp-PLA₂) is an inflammatory enzyme that catalyzes the hydrolysis of oxidized phospholipids into lysophospholipids on the surface of lipoproteins, and approximately 80% of serum Lp-PLA₂ was found on the surface of LDL particles, migrating across the vascular endothelial cell layer together with LDL^[114]. Coronary arterial lesions had a high concentrations of Lp-PLA2 in the necrotic core^[394], which suggested its involvement in atherosclerotic plaque rupture^[114]. Intracellular parasites acquire host cholesterol that endocytosed by the LDL pathway, a process that is specifically increased in *T. gondii* infected host cells^[149], and interestingly, PLA₂ was found to be implicated in *T. gondii* cell invasion through increasing their penetration^[395].

T. gondii RELEASES FROM INFECTED HOST CELLS INTO THE EXTRACELLULAR SPACE SMALL VESICLES KNOWN AS EXOSOMES WHICH CONTAIN PATHOGEN-ASSOCIATED MOLECULAR PATTERNS (PAMP) AND OTHER BIOMOLECULES THAT STIMULATE PROINFLAMMATORY RESPONSES, AND MAY THEREFORE PLAY A ROLE IN THE PATHOGENESIS OF ATHEROSCLEROSIS

Extracellular vesicles (ECVs) are carriers of PAMP and damage-associated molecular patterns (DAMP), cytokines, autoantigens, and tissue-degrading enzymes and have a possible role in pathogenesis of several inflammatory states and various clinical entities, including infections, atherosclerosis, and autoimmune diseases. ECVs contain proteins, lipids, nucleic acids from the cell from which they derived, and participate in intercellular communication^[396].

Bhatnagar et al[397] found that THP-1 macrophages infected with intracellular pathogens, including T. gondii virulent RH strain, released from cells small 50-100 nm diameter vesicles, known as exosomes, into the extracellular space. These exosomes, when exposed to uninfected macrophages, stimulated a proinflammatory TNF-α response in a TLR2 or TLR4 receptor- and MyD88-dependent manner, and exosomes, not apoptotic vesicles, were responsible for the proinflammatory activity^[397]. Exosomes are released, especially during cellular stress, from many different cell types^[398], and dendritic cells and B cells expressed MHC class I and II molecules as well as costimulatory particles, and have been shown to promote T-cell activation[399-401]. Exosomes from a mouse dendritic cell line incubated with T. gondii-derived antigens triggered a strong systemic humoral immune response protective against infection with the pathogen^[402]. T. gondii infection of human foreskin fibroblasts also caused the production of exosome-like vesicles containing a unique array of mRNA and miRNA transcripts described in the literature data as having abnormal neurologic effects^[403-406]. The vesicles from infected cells could attach and deliver their contents to uninfected cells, and their RNA contents are not just reflecting a cytoplasmic RNA, thereby causing the function of these cells to be altered $^{[403]}$.

Several stimuli, such as proinflammatory cytokines, lead to ECVs release from monocytes and macrophages in the atherosclerotic lesion[407]. ECVs derived from circulating in blood vascular endothelial cells, VSMCs, macrophages, and other circulating immune cells mainly have proinflammatory properties^[408], and for example proinflammatory cytokines and NO can trigger blood endothelial cells-derived ECVs production^[409]. Moreover, recent findings have associated lymphatic dysfunction with the pathogenesis of atherosclerosis, hypercholesterolemia, and cardiovascular diseases^[410]. Inflammatory cell accumulation in peripheral tissues drives several chronic inflammatory diseases, including atherosclerosis and obesity[261] in which lymphatic transport has been found defective[410-412]. Milasan et al[413] showed that diverse ECs inclusively derived from erythrocytes and platelets were present in lymph of healthy animals, and their levels were modulated in lymph during atherosclerosis. It must be noted that circulating red blood cells may be infected with T. gondii^[414], human and chicken thrombocytes have phagocytic capacity[415,416], and adherence of platelets to tachyzoites of the parasite together with disruption of surface membranes and cytoplasmic contents of the organisms, were observed ultrastructurally^[417]. Although chicken erythrocytes and platelets were found to be infected with the parasite, the pathogen was however not capable of replication in these cells[418].

In summary, the parasite which infect different cell types, participates in the production of many ECVs. Circulating ECVs may contain remnants of *T. gondii* tachyzoites, such as PAMP, DAMP and other microbioparticles that are dispersed to various organs, and may take part in triggering persistent inflammation, lymphatic drainage dysfunction, and development of atherosclerosis and various cardiovascular events.

ANEMIA, HEMORRHAGE, FIBRIN DEPOSITION IN TISSUES, AND DISTURBANCES IN BLOOD COAGULATION OBSERVED DURING T. gondii Infection May Well EXPLAIN SIMILAR ABNORMALITIES CHARACTERISTIC FOR ATHEROSCLEROSIS

In mammals, T. gondii infects all nucleated cells^[419], and thus has the potential to directly damage vascular cells, thereby causing hemorrhage and activating a pathophysiological blood coagulation response. Studies of host defense against T. gondii infection showed that protection was mediated by a robust proinflammatory $T_{\rm H}1$ type of immune response characterized by the prolific production of large quantities of IFN- γ and other proinflammatory cytokines^[32,420]. Chronic overproduction of IFN- γ induced anemia by both reducing mature erythrocyte life span in the circulation and inhibiting erythropoiesis via IL-15^[421-423]. IFN- γ also stimulated expression of genes that favored and/or stabilized fibrin deposition^[424-426]. Del Prete *et al*^[427] documented that $T_{\rm H}1$ and $T_{\rm H}2$ T-helper cells exerted opposite regulatory effects on procoagulant activity and tissue factor production by human monocytes.

Several authors reported that blood coagulation leading to fibrin deposition performed multiple protective functions during infection because fibrin: (a) restrained pathogen growth and dissemination, (b) protected against hemorrhagic pathology, and (c) probably promoted tissue repair^[428-433]. Mullarky *et al*^[434] found that in mice lacking

IFN-γ, IL-12, and/or TNF-α neither hemorrhage nor pathogen burden were responsible for fibrin levels during $\mathit{T.gondii}$ infection, and that cytokines exerted dominant regulatory roles (**Table 22**). It appeared that proinflammatory T_H1 immunity provoked hemorrhage via IFN-γ while independently promoting the deposition of fibrin via TNF- $\alpha^{[434]}$. Earlier studies showed that TFN-a may favor fibrin deposition by stimulating vascular wall permeability^[435-437], and the cytokine promoted the formation of fibrin as well as fibrin-associated adhesions and abscesses in a model of septic peritonitis^[432]. TNF- α also increased expression of procoagulant biomolecules^[438,439], including PAI generation^[24-26], decreased expression of anticoagulant factors^[438-440], and/or suppressed fibrin degradation in human endothelial cells^[434,441-443].

Blood coagulation disturbances leading to fibrin deposition frequently accompanied T_H1 type of immune responses, including autoimmunity[424,429,444-446]. Studies using fibrinogen-deficient mice revealed pathological roles for fibrin (ogen) during atherosclerosis^[447], and reported that fibrin can upregulate expression of proinflammatory cytokines and chemokines in humans $^{[448]}$. Johnson et $al^{[429]}$ demonstrated that blood coagulation performed important protective function during T. gondii infection by suppressing hemorrhage evoked by IFN-γ, a critical proinflammatory mediator of T_H1 immunity. Fibrin deposition mediated protection of host tissue against T. gondii infection-stimulated immunopathology^[429]. These investigators[429] established that increased production of fibrin during T. gondii infection prevented infection-stimulated blood loss, thereby performing a protective function evoked by IFN-y, a critical immune mediator elicited by the parasite during host robust T_H1 type immune response. Since IFN-y and other cytokines stimulated also expression of genes that promote and/or stabilize fibrin deposition[424-426], it may suggest that the enhanced T_H1 immune response evolved means to locally up-regulate blood coagulation activity, thereby protecting against its own destructive bioprocesses^[429].

Plasminogen activator inhibitor-1 (PAI-1)

PAI-1, a prothrombotic protein and the fast-acting inhibitor of plasminogen activators, is the principal regulator of endogenous fibrinolytic enzyme system. Disruption of the PAI-1 gene was associated with increased resistance to thrombosis and with a mild hyperfibrinolytic state characterized by enhanced in vivo clot lysis^[449,450]. Low fibrinolytic capacity has been associated with an increased risk of recurrent major cardiovascular disorders^[451]. Nilsson et al^[36] showed that unsaturated fatty acids, i.e. oleic, linoleic, and linolenic acid, and eicopentaenoic acid, caused a significant increase in PAI-1 secretion and mRNA levels from cultured human umbilical endothelial and EA.hy926 type cells. Transfection experiments demonstrated that addition of linolenic acid and eicosapentaenoic acid markedly increased PAI-1 transcription[450]. It was found that unsaturated fatty acids induced the same complex as did VLDL, whereas saturated fatty acids had no effect. VLDL has been shown to induce a concentration-dependent increase in the PAI-1 secretion from human umbilical venous endothelial cells $^{[452\text{-}454]}$ and HepG2 cells in vitro[453,455]. In vivo, several studies demonstrated a strong positive correlation between the plasma VLDL triglyceride and PAI-1 activity levels^[456-458], and a VLDL response element was identified in the promoter region of the PAI-1 gene locus[459]. Addition of a triglyceride-rich emulsion also resulted in an enhanced secretion of PAI-1 by HepG2 cells[460]. These in vitro data are in line with the finding that administration of n-3 fatty acids in vivo resulted in increased plasma PAI-1 activity[461,462].

The increased level of PAI-1 may critically affect fibrinolysis

Table 22 Dissociation of roles for hemorrhage, pathogen burden, and cytokines in the regulation of fibrin deposition during acute T. *gondii* infection in mice (acc. to Mullarky *et al*^[434], with own modification).

Mouse ^a	Fibrin	Hemorrhage b	Parasites	IFN-γ	TNF-α		
IFN-γ-deficient	++++	-	++++	-	++		
IL-12-deficient	++	-	++++	+	+		
TNF-α-deficient	+/-	++	+++	++	-		
IFN-γ/TNF-α deficient	++++ ^c	-	++++	-	-		

WT, wild type mice; ^a Data were compiled from multiple experiments. A semiquantitative system scoring was used: ++++, more than 10-fold greater than levels detected in WT mice; +++, 2- to 10-fold greater than levels detected in WT mice; ++, equivalent to levels detected in WT mice; +, below levels detected in WT mice but above the assay's detection limit; -, not detected. ^b Hemorrhage denotes evidence for both hematocrit reductions (i.e. anemia) and positive tests for fecal occult blood (i.e. bleeding). (It was demonstrated that *T. gondii*-infected mice exhibited anemia due to a decrease in both survival time of red blood cells in the circulation and erythropoiesis (p < 0.02)^[423]). Fibrin levels in IFN- γ / TNF- α deficient mice were greatly elevated in comparison with WT mice yet significantly lower than those observed in IFN- γ -deficient animals.

because the stable complex formation rate of this inhibitor with tissue plasminogen activator (t-PA), urokinase-type plasminogen-activator (u-PA), and two-chain t-PA is described as $K = 10^7 M^{-1} s^{-1}$, which for PAI-1 is a few orders of magnitude higher than that for other inhibitors, such as PAI-2, protease nexin, $\alpha 2$ -macroglobulin, C'1-esterase, and $\alpha 2$ -antiplasmin ($K = 10^0 - 10^5 M^{-1} s^{-1}$). Moreover, the plasma t-PA concentration is 70 pmol/L and its biological half-life is a few minutes, whereas the respective values for PAI-1 are 1 nmol/L and 2 to 3 hrs (*in vitro*, 37°C)^[463-465].

Mullarky *et al*^[434] demonstrated an important role for the signal transducer and activator of transducer-1 (STAT1) in the suppression of infection-stimulated fibrin deposition via a STAT1-dependent pathway operating in endothelial cells, and interestingly, expression of PAI-1 also is STAT1 regulated^[466]. IFN-γ suppresses expression of PAI-1 in endothelial cells^[467], and as PAI-1 antagonizes fibrin degradation, thereby promoting fibrin deposition, STAT1-mediated suppression of PAI-1 expression in endothelial cells. This may well account for the effects of IFN-γ on fibrin deposition during infection^[434]. In addition, IL-4 (an antiinflammatory cytokine) suppresses PAI-2 formation in stimulated human monocytes^[427].

Transforming growth factor-β (TGF-β)

This interleukin belongs to the family of related cytokines that influence inflammatory processes, cell growth, differentiation, apoptosis, and gene expression $^{[123,468-470]}$. TGF- β can upregulate the expression of integrin matrix receptors and inhibitors of matrix-degrading enzymes, such as PAI- $1^{[123,468-470]}$, which plays an important role in blood coagulation and fibrinolysis. TGF- β 1-induced PAI- $1^{[123,468-470]}$ and both these major causative biofactors are important in the pathophysiology of vascular disease and tissue fibrosis. This cytokine has profibrotic properties because it induces the expression of many matrix proteins including collagens, fibronectin, tenascin, and thrombospondin, and the net effect is the increased synthesis and assembly of matrix that result in fibrotic changes in affected tissues $1^{[123]}$. Activation of TGF- β also involves generation of ROS and RNI that participate in development of atherosclerosis $1^{[123,472]}$.

Nitric oxide (NO)

Vascular endothelium-derived NO, a major mediator of endothelium-dependent vasodilation, has important antiinflammatory and antithrombotic properties, such as inhibiting leukocyte adhesion,

limiting platelet adhesion and aggregation, and the expression of PAI-1[473-475]

Factor XIII (FXIII)

Catani et al[476] demonstrated that blood coagulation FXIII (fibrin stabilizing factor, plasma $t_{1/2} = 3-5$ days) was a target for NO both in vitro and in vivo. NO donors inhibited FXIII activity in a dosedependent manner and inhibition occurred by S-nitrosylation of a highly reactive cystein residue. NO exerts its biological effect by interacting with a variety of molecular targets, mainly using metals such as iron, tiols and oxygen as reactive groups. NO has also been shown to interact with heme- and non-heme iron-containing proteins, thus regulating their activities. Nitrosonium ion (NO+) can react with protein thiol groups causing S-nitrosylation, thus affecting the function of these proteins. Interestingly, hydrogen sulfide (H2S), an important gaseous signaling molecule that serves many important regulatory roles in pathophysiological states through NO, inhibited atherosclerosis through upregulating protein S-nitrosylation level in aorta VSMCs^[477]. H₂S significantly reduced the aortic atherosclerotic lesion area and inhibited lipid and macrophage accumulation and vascular smooth muscle cell proliferation in apoE-/- mice[477].

von Willebrand factor (vWF)

Elevated vWF levels are strongly associated with an increased risk of ischemic cardiovascular events. vWF probably participate in the process of atherogenesis, because results from animals studies suggested that at arterial branch point predilection sites, vWF deficiency or blockage had a protective effect against atherosclerosis^[478]. Endothelial secretion of vWF from intracellular organelles is required for platelet adhesion to the injured vessel wall, because vWF binds platelets with collagen fibers in subendothelial layer and serves as a basis of platelets adhesion and aggregation. Torisu *et al*^[479] showed that autophagy regulated endothelial cell processing, maturation and secretion of vWF (autophagy is a reparative process by which cytoplasmic components are sequestrated in double-membrane vesicles and degraded on fusion with lysosomal compartments).

Effect of blood flow rate

Since the blood flow rate is the main determinant of in *vivo* platelet-vessel wall interaction^[480], the platelets would tend to accumulate at the points of irregularities in blood flow dynamics, and might then release vasoactive agent(s) (possibly serotonin) and PAI-1^[481], which is also liberated from endothelial cells of vessel walls during venous stasis^[482]. It should be noted that the platelet pool of PAI-1 is four- to six-fold greater than that of plasma^[481,483].

Serotonin (5-HT)

5-HT, a potent vasoconstrictor in the large cerebral arteries, is considered to play a key role in atherosclerosis and to be implicated in ischemic cerebrovascular events followed by delayed neuronal death^[484]. 5-HT released from activated platelets, not only accelarated aggregation of platelets but also promoted mitosis, migration, and contraction of VSMCs, and these effects contributed to thrombus formation and atherosclerosis^[484]. Serotonin stimulated significantly the expression of PAI-1 and tissue factor mRNA in a concentrationand time-dependent manner through 5-HT_{2A} receptors, thus increasing procoagulant activity and reducing fibrinolytic activities of endothelial cells^[485]. 5-HT-stimulated endothelial cells secreted a T lymphocyte-specific chemotactic cytokine with competence growth factor activity^[486]. 5-HT alone and combined with thromboxane A₂,

potently induced vascular smooth muscle cell proliferation [487,488]. However, there is still controversy regarding the effects of 5-HT on endothelial cell proliferation [489]. Finally, Suguro *et al* [490] found that 5-HT increased ACAT-1 activity in a concentration-dependent manner in primary monocyte culture, and suggested that 5-HT may play an important role in macrophage-derived foam cell formation. It must be noted that hyperserotoniennal has been reported in autistic and mentally retarded children [491,492], as well as in the first degree relatives of these children [493,494], and notably, Prandota *et al* [495] found a significant association between autism and *T. gondii* infection ($\chi^2 = 8.11$, p < 0.0043; OR = 7.2; 95% CI: 1.56-33.7).

Glycated LDL

Artwohl *et al*^[496] found that diabetic LDL triggered apoptosis in vascular endothelial cells. It was reported that glycated LDL increased adhesion molecule expression^[497] and modulated the fibrinolytic potential of vascular endothelial cells^[498]. Furthermore, both *in vitro*- and *in vivo*-glycated LDL altered vasoactive response by attenuating NO synthesis^[499]. The toxicity of glycated LDL and its role in the pathogenesis of atherosclerosis appeared to be related to its prolonged presence in the circulation and impaired cellular uptake^[498,500]. It should be noted that *T. gondii* infection acquired prenatally and/or after birth may be responsible for development of both type 1 and 2 diabetes mellitus^[501].

Magnesium deficiency

Maier *et al*^[502] demonstrated that low magnesium concentrations reversibly inhibited endothelial cell proliferation, probably due to an up-regulation of IL-1 synthesis, a proinflammatory cytokine and potent inhibitor of endothelial cell growth. They observed also the up-regulation of VCAM-1 and PAI-1 levels after Mg deficiency.

All together, overproduction of proinflammatory cytokines, such as IFN- γ , TNF- α , and NO, during latent chronic *T. gondii* infection may lead to serious physiopathological disturbances in the host, including anemia, hemorrhage, and blood coagulation disorders, i.e. the abnormalities that also play an important role in the clinical course of atherosclerosis.

IMPORTANT ROLE OF IRON IN DEVELOP-MENT OF ATHEROSCLEROSIS AND NEURO-DEGENERATIVE DISEASES

Atherosclerosis

According to the "iron hypothesis", iron overload is believed to be detrimental for the cardiovascular system, thus promoting atherosclerosis development and progression^[503,504]. Excess of iron generates oxidative stress that hallmarks neurodegenerative diseases and atherosclerosis, a frequent disorder of aging^[505]. Patients with Alzheimer's disease and Parkinson's disease showed a dramatic increases in their brain iron content which may contribute to the pathogenesis of these disorders, but so far the reason for iron accumulation is unknown^[506].

Iron accumulates in human atherosclerotic lesions^[507], and can enter into the lesion in the form of free hemoglobin (Hb) that is released upon intravascular hemolysis or intraplaque hemorhage^[508]. Cell-free Hb when oxidized releases heme and induces oxidative modification of LDL^[509]. Iron mediates the oxidative modification of LDL, an important contributing factor to the pathogenesis of atherosclerosis^[510]. Cholesterol levels in atherosclerotic lesions correlated with iron deposits^[511], and within the plaque, iron deposition and ferritin induction have been observed in endothelial

cells and macrophages in early human lesions, and additionally in VSMCs in late lesions^[503]. Interestingly, in hypercholesterolemic pigs, Gerrity^[512] found that monocytes were adherent to the lesion areas of aortic endothelium, in endothelial junctions, and in the intima. Intimal monocytes and altered intimal monocytes with enlarged cytoplasm and containing few lipid droplets were found to be phagocytic by their uptake of ferritin, which had penetrated the intima after intravenous injection. Circulating monocytes and those adherent to the endothelial surface did not contain ferritin in the animals^[512].

Hb derived iron is an important factor in determining macrophage differentiation and function in areas of intraplaque hemorrhage within human atherosclerosis. Habib et al[160] showed that macrophages in areas of intraplaque hemorrhage demonstrated reduced intracellular iron and ROS which triggers production of antiinflammatory cytokines as well as genes involved in cholesterol efflux. Iron is a powerful catalyst in the generation of toxic free hydroxyl radicals through the oxidation of its ferrous (Fe2+) (very toxic) to ferric (Fe³⁺) via the Fenton reaction (Fe²⁺ + H_2O_2 -> Fe³⁺OH⁻ + OH)^[513], which are deleterious to living cells and most macromolecules^[514]. Hydroxyl radicals can depolymerize polysaccharides, cause DNA breaks, inactivate enzymes, and initiate lipid peroxidation $^{[160,515\text{-}517]}.$ Extracellular ferric anion is bound to human serum lactoferrin (Lf) and transferrin, and transferrin is the iron transporter that allows cellular iron uptake^[518]. Lf and transferrin maintain Fe³⁺ in a soluble and stable oxidation state, avoiding the generation of toxic free hydroxyl radicals[514].

Hemolysis is often a result between the interaction of erythrocytes and mature atheromas resulting in the transition of ferrous to ferric forms of Hb which additionally leads to lipid peroxidation^[519], triggering endothelial activation, smooth muscle cell proliferation and macrophage activation^[503]. Oxidized form Hb can also act as a proinflammatory agonist that targets vascular endothelial cells^[520].

Hydrogen peroxide (H_2O_2) generated by mitochondria and other organelles permates in the lumen of secondary lysosomes which contain iron derived from cellular structures undergoing autophagic degradation^[521,522]. The interaction between reactive ferrous iron and H_2O_2 resulted in the generation of hydroxyl radicals inducing lipid peroxidation and ceroid formation^[523]. Lee *et al*, ^[524] reported that iron and ceroid deposits colocalized either extracellularly or intracellularly in foam cell-like macrophages or SMCs of advanced atherosclerotic plaques. Many cells in advanced human plaques contained a large number of ceroid-containing lysosomes, and impaired autophagy stimulates further accumulation of damaged mitochondria, increased ROS generation, additional accumulation of iron and enhanced non-degradable oxidation products such as ceroid^[521].

T. gondii infection

The parasite must acquire nutrients from its host to survive and ensure its proliferation^[525]. *T. gondii* is an auxotroph for iron^[526,527], cholesterol^[149,528], tryptophan^[529], arginine^[526,530], polyamines^[531], purines^[532], and other essential nutrients^[146]. Parasites have an ability to acquire growth-essential iron from their hosts^[533]. Possible sources of intracellular host iron include heme, iron released from transferrin at pH 5.5-6, and ferritin^[533]. The metal is needed for catalysis of DNA synthesis and for a variety of enzymes concerned with electron transport and energy metabolism. Iron may also focus influx of neutrophils to sites of infection by generating oxidant-sensitive mediators^[534].

Dimier and Bout $^{[527]}$ demonstrated that T. gondii invaded and proliferated in cultured primary rat enterocytes. Intestinal epithelial cells represent a rapidly renewing cell population, and their cell

turnover has a mean duration time of 2-3 days *in vivo* and a few hours *in vitro*^[535]. Experiments that used Fe²⁺ salt as well as carrier and chelator suggested that IFN- γ -treated enterocytes inhibited *T. gondii* replication within these cells by limiting the availability of intracellular iron to the parasite^[527,536]. Data from an *in vitro* model of human monocytes infected with other pathogens, such as *Legionella pneumophila*^[537], *Histoplasma capsulatum*^[538], or *Trypanosoma cruzi*^[539] also have shown that limiting the availability of iron may represent a broadly effective intracellular antimicrobial mechanism. IFN- γ inhibited the replication of *T. gondii* also in endothelial cells^[56], fibroblasts^[540], and retinal pigment epithelial cells^[57]. In fibroblasts and retinal pigment epithelial cells, the mechanism of inhibition was via tryptophan starvation, while in astrocytes, the mechanism of IFN- γ inhibition was independent of iron deprivation and reactive oxygen intermediates^[541].

Several pivotal proinflammatory cytokines such as IFN-γ, TNF-α, IL-1, and IL-6 (these molecules are profusely generated during chronic T. gondii infection) participated in iron homeostasis, and the sequestration of this metal, mediated by these biomediators, was considered protective^[542]. Proinflammatory cytokines and the acute phase protein hepcidin affected iron homeostasis leading to the accumulation of the metal ion in macrophages and subsequently to anemia of inflammation^[543]. Hepcidin inhibited iron efflux through binding to and induction of the degradation of ferroportin1, a multipass protein found in the basolateral membrane of enterocytes^[544]. Iron and the hormone erythropoietin modified innate immune responses by influencing IFN-γ-mediated (iron) or NF-κB inducible (erythropoietin) immune effector pathways in macrophages. Thus, the macrophages loaded with iron lost their ability to kill intracellular pathogens via IFN-γ-mediated effector pathways, such as NO generation[543]. On the other hand, NO directly controlled intracellular iron metabolism by activating iron regulatory protein, a cytoplasmic protein that regulates ferritin translation^[543]. It was reported that NO mediated non-heme iron release from ferritin[545], and ferritin, a hepatic protein and cytoprotective antioxidant of endothelium, may counteract some redox activity via ferroperoxidase in the vasculature^[546]. Iron also regulated NOS activity by controling nuclear transcription of cytokine-inducible NOS mRNA[541]. Weiss et al[547] showed that increased intracellular iron levels led to a decrease of NOS activity, whereas depletion of intracellular iron strongly enhanced the enzyme activity in IFN-γ/LPS-stimulated macrophages, i.e. alterations (increases as well as decreases) in cellular iron availability caused regulation of NOS transcription in J774 cells^[539].

Macrophages and other host cells activated by IFN- γ can be induced to form a flavoprotein that converts L-arginine to NO + L-citrulline^[548]. NO causes efflux of non-heme iron from infected host cells. In the absence of L-arginine, IFN- γ -induced infected cells can lower their net uptake of iron. Cellular depletion of the metal via either mechanism suppresses DNA synthesis as well as the functioning of aerobic respiratory enzymes. Macrophage regulation of growth of other host cells during embryogenesis, immune responses, or immunosurveillance might involve iron depletion^[548].

Novel atherosclerotic markers indicating neutrophil activation represent Lf, an innate immune system protein with antiinflammatory and antioxidant properties^[549,550]. Immunohistochemical staining of Lf was found in advanced human atherosclerotic carotid artery, as well as its co-localization with neutrophils and T and B lymphocytes in aortic plaques^[551]. Lf appeared to be one of nine genes specifically induced predominantly in atherosclerotic aortic plaques^[551]. Lf, a natural defense iron-binding protein is produced and secreted by glandular epithelial cells and circulating polymophonuclear neutrophils, and

found mainly in milk, saliva, mucosal and other secretory fluids, has been found to possess antiinflammatory, antibacterial, antiparasitic, antimycotic, antiviral, and antineoplastic activity[552,553]. A principal function of Lf is that of scavenging free iron in fluids and inflamed areas so as to suppress free radical-mediated damage and decrease the availability of the metal to invading microbial and neoplastic cells. Lf normally maintains an iron saturation value of 25-35%, and at values above 30%, transferrin (present mainly in serum and lymph) begins to lose its effectiveness as a scavenger of hazardous iron^[552,554]. Lf, a 78-kDa glycoprotein of the transferrin family, is structurally similar to transferrin with about 44% homology, and similar to transferrin Lf can reversibly bind two atoms of iron (Fe³⁺). T. gondii utilize iron bound to inorganic or organic chelates, such as serum Lf and transferrin and/or intracellular iron compounds (Hb, hemin, ferritin and others[555,556]. Iron uptake by the parasite was limited by IFN-γ through an unknown mechanism^[518]. Dziadek et al^[557] found in human CaCo-2 epithelial cells, and mouse L929 fibroblasts, that extracellular T. gondii tachyzoites of BK strain (genotype I) bind specifically human serum Lf but not transferrin, and the binding was concentration-dependent. Moreover, the multiplication of the parasite was inhibited by human Lf high concentrations (> 100 mg/ mL) in both cell lines^[553]. Virtually all cells express a transferrin receptor on their surface, and the quantity of receptor molecules reflects the cellular iron requirements^[518]. It should be noted that host cell expression of transferrin receptor was upregulated in T. gondii infected cells, suggesting that the parasite utilized the host endocytic pathway for iron acquisition[518,526]. Since the receptors for human Lf also are expressed on many human cells, they could provide a mechanism for the interaction between Lf binding T. gondii tachyzoites and host cells[558]. It appeared that the presence of iron was required for the Lf molecule to stimulate an increase of uptake and killing of an intracellular parasite by murine macrophages and human monocytes[559,560], especially that T. gondii binds LF and not transferrin^[557]. T. gondii membrane proteins ROP4 and ROP2 were found to be involved in human Lf acquisition[561]. Human Lf displayed cytoprotective (in a concentration-dependent manner) and antioxidant effects against H₂O₂-induced oxidative stress in human umbilical vein endothelial cells^[562]. Immunomodulatory effects of Lf include interacting with monocytes and macrophages and modulating their function during inflammatory and infectious processes, e.g., increasing cytotoxic activity, cytokine production (T_H1) and expression of surface molecules^[563], enhancing phagocytosis, induction of apoptosis, and promoting T_H2 cell responses in intestinal mucosa^[564]. Moreover, Lf displayed beneficial effects on plasma lipid levels, because circulating Lf levels were inversely associated with fasting triglycerides $(p = 0.001)^{[565]}$, and Lf administration led to decreased plasma triacylglycerol and free fatty acids levels, and to increased plasma HDL-cholesterol concentration accompanied by decreased hepatic cholesterol and triacylglycerol contents in rodents[566].

In summary, iron accumulates in human atherosclerotic lesions and play an important role in the pathogenesis of these disorders because this metal may focus influx of neutrophils to sites of infection by generating oxidant-sensitive mediators. T. gondii is an auxotroph for iron and other nutrients. Experimental data suggested that IFN γ -treated enterocytes inhibited T. gondii replication within these cells by limiting the availability of intracellular iron to the parasite. Proinflammatory cytokines and the acute phase protein hepcidin affected iron homeostasis leading to the accumulation of the metal ion in macrophages and subsequently to anemia of inflammation. Iron modified innate immune responses by influencing IFN- γ -mediated

immune effector pathways in macrophages. The macrophages loaded with iron lost their ability to kill intracellular pathogens via IFN-ymediated effector pathways, such as NO generation. NO mediated non-heme iron release from ferritin, and ferritin may counteract some redox activity in the vasculature. Extracellular T. gondii tachyzoites bound specifically human serum Lf, and the parasite membrane proteins ROP4 and ROP2 were involved in Lf acquisition. It must be noted that Lf is one of 9 genes specifically induced predominantly in atherosclerotic aortic plaques, and its principal function was that of scavenging free iron in fluids and inflamed areas and decrease the availability of the metal to invading microbial cells. Thus, a marked increase of iron content in both atherosclerotic lesions and brain tissues in neurodegenerative diseases may reflect a defense of the host against infection with the parasite and contribute to the pathogenesis of these disorders, especially that iron regulates NOS activity by controling nuclear transcription.

POSSIBLE LINK BETWEEN ELEVATED PLASMA LEVELS OF TGF-β1 IN ATHEROSCLE-ROSIS, AND T. gondii INFECTION. DUAL ROLE: TGF-β INHIBITS INFLAMMATION BUT INCREASES INTRACELLULAR PARASITE REPLICATION

Atherosclerosis

Inflammation play an important role in atherosclerosis [^{567]}. In advanced atherosclerosis, the serum concentration of active TGF- β was found to be severely depressed [^{568,569]}. Blann *et al* [^{570]} demonstrated also a significantly increased serum levels of TGF- β receptors in atherosclerotic patients with ischemic heart disease and peripheral vascular disease, which correlated with serum total cholesterol levels (p < 0.001). Several authors further established that this cytokine reduced atherosclerosis by dampening T cell activation [^{571,572]}, while disruption of TGF- β signaling in T cells resulted in acceleration of this inflammatory process [^{571,573,574]}.

TGF- β belongs to the family of related cytokines that influence cell growth, differentiation, apoptosis, inflammatory processes, and gene expression [468-470]. This is a multipotent cytokine with antiinflammatory activities, such as inhibition of proliferation, maturation and/or activation of macrophages, lymphocytes and NK cells [575-578]. TGF- β is an important regulator of inflammation, being proinflammatory at low concentrations and antiinflammatory at high concentrations [579]. Activation of TGF- β involves reactive oxygen species [472]. This cytokine is secreted by a number of cells, including regulatory T cells, macrophages, endothelial cells, dendritic cells, platelets, and vascular smooth muscle cells [580,581]. Tonkin and Haskins [582] demonstrated that regulatory T cells transfer caused a reduction in the number of effector T_H1 T cells and macrophages, and also inhibited effector T cell cytokine and chemokine production.

TGF-β has profibrotic properties because it induces the expression of many matrix proteins including collagens, fibronectin, tenascin, and thrombospondin, and the net effect is the increased synthesis and assembly of matrix that result in fibrotic changes in affected tissues^[123]. TGF-β system mediated diabetic renal hypertrophy and fibrosis build-up due to the extracellular matrix production^[583,584], the cytokine was found to induce vaginal tissue fibrosis in animal model^[585], and the beta cell hypertrophy, beta cell damage and fibrosis, with reduction in insulin secretion, was characteristic for patients with type 2 diabetes mellitus^[501,586].

TGF- β plays a critical role in the suppression of lymphocyte proliferation and differentiation therefore preventing hazardous autoimmune responses, and its immunosuppressive effects are mediated through the inhibition of TNF- α and IL-1^[587] and blocking the induction of adhesion molecules like ICAM-1 and VCAM-1^[588-590]. Filisetti and Candolfi^[373] reported that TGF- β is well known for its immunosuppressive action on leukocyte cell lines. This cytokine was found to be an antagonist of TNF α , TNF- β , IFN- γ and IL-2^[591,592]. The antiinflammatory action of TGF- β control development of immunopathological processes related to TH1 immune response in the brain^[593] and the intestines^[594].

TGF- β can also upregulate the expression of integrin matrix receptors and inhibitors of matrix-degrading enzymes, such as PAI-1^[471], which plays an important role in blood coagulation and fibrinolysis. TGF- β 1-induced PAI-1 expression was found also in VSMCs^[471], and both these major causative biofactors are important in the pathophysiology of tissue fibrosis and vascular disease.

T. gondii infection

Host protection to T. gondii infection involves T_H1 type immune response of inflammatory cells, lymphocytes and macrophages with enhanced production of IFN- γ , TNF- α , and IL-1 $\beta^{[373,595]}$. Activation of macrophages plays an important role in the host resistance against intracellular pathogens. Langermans et al[592] found that the IFN-yinduced toxoplasmastatic activity of macrophages was inhibited by TGF- β (mean fold increase = 6.3), which was also found for the IFNγ-induced production of TNF-α, RNI and PGE₂ by macrophages. It appeared that inhibition of TNF-α production was a key factor in the TGF-β-induced suppression of macrophage activation with respect to toxoplasmastatic activity and RNI production[592]. Activated macrophages by IFN-γ inhibit parasite replication through a number of potent microbicidal mechanisms such as oxidative[596] and nonoxidative^[597] mechanisms, as well as the induction by IFN-γ of enzyme indoleamine 2,3-dioxygenase that degrades tryptophan, which is required for the parasite replication^[598]. Barbosa et al^[599] showed that treatments with IL-10 or TGF-β1 induced a considerable augmentation in both T. gondii intracellular replication and invasion into BeWo trophoblastic cells, in contrast with HeLa cells. BeWo cells were unable to control replication of the parasite even in the presence of exogenous IFN- $\gamma^{[600]}$. Nagineni et al^[210] reported that in human retinal pigment epithelial cultures TGF-β also enhanced parasite replication. Soluble extracts of T. gondii stimulated secretion of both TGF-β1 and TGF-β2 significantly. T. gondii infection completely inhibited secretion of the active form of TGF-β2. In addition, Malipiero et al^[601] found that endogenous TGF-β suppressed host defense against pathogen infection also in the central nervous system.

TGF- β is downregulating tissue inflammation and immunity in the gastrointestinal tract (intraepithelial lymphocytes via the secretion of TGF- β are critical in preventing inflammation of the small intestine villi following parasite infection), the central nervous systems, and the eye, where latent *T. gondii* reside and often become reactivated [594,602,603]. TGF- β was found to increase *in vitro* replication of *T. gondii* in retinal cells[604]. Elevated expression of TGF- β in vitreous, retina and retinal pigment epithelium has been correlated closely with retinal fibrosis and choroidal neovascularization[210]. One cannot therefore exclude that the development of fibrotic changes in atherosclerotic lesions are associated with the increased levels of this cytokine due to *T. gondii* infection because TGF- β belongs to the biomediators favouring growth of the parasite [605], and at the same time mediates balance between inflammation and fibrosis during plaque formation [6066].

Activated macrophages control growth by NO production [607].

However, *T. gondii* active invasion inhibits NO production, allowing parasite persistence. The mechanism used by *T. gondii* to inhibit NO production persisting in activated macrophages depends on phosphatidylserine exposure. TGF- β 1 led to iNOS degradation, actin filament (F-actin) depolymerization, and lack of NF- κ B in the nucleus^[607].

Taken together, it seems that the increased levels of plasma TGF- β reported in patients with atherosclerosis exert both beneficial and harmful effects because although this cytokine inhibits inflammation and autoimmunity, at the same time increases *T. gondii* replication in the host cells with further development of unwanted pathophysiological consequences, including fibrotic changes in atherosclerotic lesions. Final clinical effect of TGF- β depends mainly of its serum levels, being proinflammatory at low concentrations and antiinflammatory at high levels.

MAST CELLS (MCs) PLAY AN IMPORTANT ROLE IN THE PATHOGENESIS OF ATHERO-SCLEROSIS, AND SIMULTANEOUSLY EXERT A BENEFICIAL EFFECT IN HOST DEFENSE AGAINST T. gondii INFECTION

Atherosclerosis

MCs are important in the development of both innate and acquired immunity, inflammation, and atherosclerosis [608-613]. MCs, macrophages, and neutrophils release TNF-α, IFN-γ, and IL-6 that induce expression of adhesion molecules in vascular endothelium and recruit of leukocytes, which is essential to the pathogenesis of vascular inflammatory diseases^[614]. MCs by virtue of mediator secretion and degranulation, can regulate the function of endothelial cells, T lymphocytes, macrophages, and VSMCs^[613]. The cells produce a number of biomediators, including histamine, tryptase, chymase, cathepsin G, platelet activating factor, leukotrienes, thromboxanes, prostaglandins, matrix-degrading metalloproteinases (MMP-1, MMP-3), inflammatory cytokines (TNF-α, IL-4, IL-5, IL-6, IL-13), and chemokines (MCP-1, IL-8, RANTES)[613]. MCs can assist in the recruitment of monocytes and lymphocytes into vascular tissue, thereby propagating the proinflammatory response^[614], and are present in atherosclerotic lesions and thrombotic sites^[615,616]. MCs were found in close association with macrophages and extracellular lipids, and at the sites of foam cell formation in atherosclerotic lesions^[616]. MCs facilitate foam cell formation by promoting the formation of LDL aggregates, native and oxLDL uptake by macrophages, and by interfering with macrophage cholesterol removal by HDL[615-622]. Moreover, the cells provide a general anticoagulant function through actions of tryptase (fibrinogenolysis, inactivation of high molecular weight kininogen), heparin proteoglycans (anticoagulant, inhibition of platelet aggregation), and therefore might slow thrombus formation at the sites of atheromatous plaque rupture[613,620].

T. gondii infection

The parasite is a potent T_H1 type cytokine inducer, and while these proinflammatory cytokines are required to survive infection, their overproduction may lead to pathology and death of the host. Effector cells against *T. gondii* include monocytes^[617], IFN-γ-activated macrophages^[420,624], neutrophils^[623], and platelets^[417]. MCs also participate in host defense against the parasite. **Table 23** presented effects of mast cells incubation with *T. gondii* tachyzoites and influence of two different selective leukotriene inhibitors (5-lipooxygenase inhibitor A-63162 and 5-lipoxygenase-activated

protein inhibitor MK-886) on MC degranulation and *T. gondii* cytotoxicity^[625].

MCs cells play a crucial role in the development of a protective immune response during resistance against *T. gondii* oral infection because of their ability to release proinflammatory mediators^[625-629]. The cells degranulate and release LTB4, which damage tachyzoites^[625]. Earlier study showed that LTB4-induced *T. gondii* cytotoxicity was characterized by vesiculation of the surface membrane and loss of the cytoplasmic contents of the organisms^[624], and the same cytotoxic changes were observed in tachyzoites after incubation with MCs^[625]. Although the parasite inhibited mast cell degranulation by suppressing the mobilization of intracellular Ca²⁺ mediated by C-phospholipase^[630], many biomediators, such as IL-6, can also be released from mast cells "selectively", without degranulation^[631,632]. In addition, it appeared that reactive oxygen species were not implicated in the mast cell-mediated toxoplasmacidal activity^[625].

Taken together, MCs play an important role in both development of atherosclerosis and host defense against *T. gondii* infection.

PLATELETS AND SEROTONIN (5-HT) ENHANCE MACROPHAGE FOAM CELL CELL FORMATION AND ARE CYTOTOXIC TO T. gondii TACHYZOITES (IN THE ABSENCE OF ANTIBODY)

Atherosclerosis

5-HT, a potent vasoconstrictor in the large cerebral arteries, is considered to play a key role in atherosclerosis and to be implicated in ischemic cerebrovascular events followed by delayed neuronal death^[484]. Ban *et al*^[633] found that plasma 5-HT concentrations, serum levels of hepatocyte growth factor (HGF), LDL cholesterol, and hsCRP, carotid artery intima-media thickness (IMT) and plaque frequency were significantly greater in patients with vascular dementia than in controls. There was a significant positive correlation of max IMT with 5-HT or HGF levels, and the authors suggested that increased plasma 5-HT levels and carotid atherosclerotic plaques may be involved in the pathogenesis and progression of vascular dementia^[633].

Atherogenesis involves platelet activation with subsequent serotonin release. 5-HT stimulated monocyte adhesion^[484,634], and enhanced macrophage foam cell formation with increased uptake of oxLDL (protein concentration 10-75 mg/L)^[188,635]. Suguro *et al*^[188] found that during differentiation of cultured human monocyte into macrophages expression of ACAT1 protein increased in a time-dependent manner. Serotonin increased ACAT1 activity in a concentration-dependent manner also in primary monocyte culture. In addition, 5-HT may contribute to inflammatory activation of the vessels during atherogenesis because it increased synthesis of IL-6 in human VSMCs, and this cytokine is a key molecule in chronic inflammation and was implicated in the progression of atherosclerosis^[636].

5-HT released from activated platelets, not only accelerated aggregation of platelets but also promoted mitosis, migration, and contraction of VSMCs, and these effects contributed to thrombus formation and atherosclerosis^[484]. Serotonin stimulated significantly the expression of PAI-1 and tissue factor mRNA in a concentrationand time-dependent manner through 5-HT_{2A} receptors, thus increasing procoagulant activity and reducing fibrinolytic activities of endothelial cells^[465]. 5-HT-stimulated endothelial cells secreted a T lymphocyte-specific chemotactic cytokine with competence

Table 23 Ultrastructural morphometric changes of MCs after interaction with *T. gondii* tachyzoites and effects of the selective inhibitors on these alterations (acc. to Henderson *et al*^[625]; with own modifications).

Reaction mixture	Mast cell degranulation, degranulated cells/ total cells (%)	T. gondii cytotoxic changes, damaged T. gondii/total T. gondii (%)
Mast cells	27/305 (8.9)	-
T. gondii	-	42/580 (7.2)
Mast cells + T. gondii	278/398 (69.8)	134/306 (43.8)
Mast cells + <i>T. gondii</i> + A-63162	71/341 (20.8) ^a	32/234 (13.4) ^a
Mast cells + T. gondii + MK-886	44/206 (21.4) ^a	36/300 (12.0) ^a
Mast cells + <i>T. gondii</i> + indomethacin	252/399 (63.2)	144/389 (37.0)

Mast cells (2 x 10⁶) and *T. gondii* RH strain (1.2 x 10⁷) were incubated in absence or presence of A-63162 (10⁶ M), MK-886 (10⁶ M), or indomethacin (10⁻⁵ M) in 1 mL of Tyrode's buffer for 30 min at 37°C. Percent of degranulated mast cells and damaged tachyzoites was determined by morphometry. Data are means of 2-3 experiments for each condition. ^a p < 0.05 vs mast cells + T. gondii.

growth factor activity^[466]. This biomediator, alone and combined with thromboxane A₂, potently induced VSMC proliferation^[467,468], but there is still controversy regarding its effects on endothelial cell proliferation^[469].

T. gondii infection

Platelets may play an important role in the host defense against T. gondii. Human platelets were found to be cytotoxic to the parasite tachyzoites^[637]. Adherence of platelets to the pahogen and disruption of surface membranes and internal architecture were observed ultrastructurally. The interaction of T. gondii with platelets resulted in a marked increase in TXB2 generation compared with that by unstimulated platelets^[631]. Moreover, Henderson et al^[638] demonstrated that human platelets incubated with T. gondii tachyzoites released oxygenated products of both arachidonic acid and linoleic acid, 13-hydroxyoctadecadienoic acid (13-HODE) (87.7%) and 9-HODE (12.3%). Further study[639] revealed that 13-HODE, a product of linoleic acid metabolism, at concentrations higher or equal to 10(-8) M rapidly induced cytotoxic changes in the parasite, including leakage of cytoplasmic contents in the space between the inner and outer parasite bilayer membrane units, which was followed by intracellular vacuolation and loss of cytoplasmic contents. Interestingly, Shamseddin et al^[640] showed that conjugated linoleic acid, a natural component of milk and diary products, in concentrations 50-250 µM induced apoptosis in virulent RH and avirulent Teheran strains of T. gondii, in HeLa cells.

T. gondii INFECTION INCREASES LEPTIN LEVELS WHICH PLAY AN IMPORTANT ROLE IN TRIGGERING INFLAMMATION AND PARTICIPATE IN DEVELOPMENT OF ATHEROSCLEROSIS

Atherosclerosis

Leptin, the adipocyte-secreted hormone, is mainly produced by white adipose tissue and circulates in the blood in levels proportional to the fat mass, and both central and peripheral administration of leptin inhibited apetite and adiposity^[641,642]. This adipocytokine modulates immune responses^[643], metabolism and inflammation^[644], and is a potential linker between obesity and

chronic inflammation^[645]. Structurally, it resembles IL-2 in particular and is a crucial T-cell growth factor^[646], and a potent negative regulator of IL-2 induced NK cell cytotoxicity^[647]. The levels of this biomediator were increased in infections and autoimmune disorders, or after exposure to inflammatory stimuli such as LPS, TNF- α , and IL-1^[641,648].

Leptin accelerated CEs accumulation in human monocyte-derived macrophages by increasing ACAT1 expression and cholesterol efflux from these cells^[208]. In human atherosclerotic lesions, leptin mediated atherosclerosis *in vitro* and *in vivo*, and induced (at doses 10-40 ng/mL) proliferation of the human umbilical vein endothelial cells and elevation of matrix metalloproteinases (MMPs) MMP-2 and MMP-9 expression, in a dose-dependent manner^[649]. Leptin caused also concentration- and time-dependent increases in IL-6 production^[390], which is the predominant mediator of the acute phase response, an innate immune mechanism triggered by infection and inflammation^[388,389]. IL-6 levels were elevated in chronic inflammatory conditions, such as rheumatoid arthritis^[650,651].

T. gondii infection

It was demonstrated that increased leptin levels correlated with enhanced concentrations of inflammatory markers in morbidly obese individuals^[645], and recently, Reeves *et al*^[652] reported a significant positive association between T gondii seropositivity and obesity (p=0.01). Individuals who were obese had a markedly higher T gondii IgG titers compared to normal weight participants^[652]. Also Baltaci and Mogulkoc^[653] found that rats infected with T. gondii had significantly increased plasma leptin levels four weeks after the intraperitoneal injection of live parasites as compared with control animals (**Table 24**).

It must be emphasized that in obese prepubertal children statistically significant positive correlations were found between leptin and IL-2, IL-1 β , IL-6 or TNF- α serum concentrations^[654] (**Table 25**), and interestingly, psychotropic drugs that induce weight gain in psychiatric patients also clearly activated the TNF- α system^[655].

Multiple cytokines and acute inflammation that raise leptin levels may have a potential role also in inflammatory anorexia [⁶⁴¹], especially that leptin treatment increased energy expenditure (oxygen consumption) in animals [⁶⁵⁶]. It should be noted that in mice with *T. gondii* infection the dual anorexia and hypermetabolic states were associated with the elevation of TNF- α , IL-1 β , IL-2, IL-4, IL-5, IL-6, IL-10, and IFN- γ levels [^{657,658}], and it was reported that various cachectic-anorexic conditions resulted from interactions among cytokines and neurotransmitters as mediators of neurologic and neuropsychiatric manifestations of disease [^{659,660}].

In summary, *T. gondii* infection caused increased leptin levels, which were found to be correlated with the enhanced concentrations of inflammatory markers in morbidly obese individuals, and recently, a significant positive association between *T gondii* seropositivity and obesity has been reported. Because latent chronic *T. gondii* infection has a high prevalence worldwide, and 23% of the European adult population are now obese, while approximately 57% of the world's adult population is projected to be obese or overweight by 2030^[661], atherosclerosis associated with inflammation and obesity represent a serious global health hazard.

IMPORTANT ROLE OF CYSTEINE CATHEP-SINS (CAT) IN FOAM CELL FORMATION AND GENERATION OF AMYLOID IN ATH-

EROSCLEROTIC ARTERIES. INFECTION OF THE HOST VASCULAR ENDOTHELIAL CELLS BY T. gondii TACHYZOITES BECOMES AN ADDITIONAL SOURCE OF SIMILAR ENZYMES BECAUSE THE PARASITE EXPRESSES FIVE MEMBERS OF THE CATHEPSIN PROTEASES, SUCH AS ONE CATL-LIKE, ONE CATB-LIKE, AND THREE CATC-LIKE PROTEASES

Atherosclerosis

Cathepsins contribute to several pathophysiological processes, including antigen presentation in the immune system^[662,663], neuropeptide and hormone processing^[664], and collagen turnover in bone and cartilage^[665]. These enzymes have strong elastolytic and collagenolytic properties that form a distinct subgroup of atherosclerosis-related proteases because they affect extracellular matrix degradation and modulate inflammation, immune responses, and cellular functions^[666-670]. Increased expression and translocation of lysosomal cathepsins contribute to macrophage apoptosis in atherogenesis^[669].

Cathepsins of the cysteine protease family, such as CatB, C, H, F, K, L, O, S, V, W, and $X/Z^{[671]}$ function in terminal protein degradation optimally whithin acidic lysosomes^[672], and play an important role in development of atherosclerosis-based cardiovascular diseases^[673-678]. Cathepsins are expressed in macrophages, endothelial and vascular smooth muscle cells (VSMCs) of atherosclerotic lesions^[677]. Studies on vascular cells have shown that atherosclerosis-associated proinflammatory cytokines enhanced cysteine cathepsin expression and activity. It was demonstrated that although VSMCs and macrophages physiologically express negligible levels of CatL, S, and K, incubation of these cells with proinflammatory cytokines, such as TNF- α , IL-1 β , and IFN- γ , significantly induced the expression and secretion of those cathepsins and their collagenolytic and elastolytic activities^[673,676,679].

Table 24 Body weight and plasma leptin concentrations in rats intraperitoneally infected with *T. gondii* (acc. to Baltaci & Mogulkoc[653], with own modification)

with own mountains.								
Study group	Body weight before the study (g)	Body weight after four weeks (g)	Plasma leptin levels (ng/mL)					
Control rats $(n = 10)$	266.00 ± 32.81	270.50 ± 33.70	4.09 ± 1.15					
Infected animals (n = 10)	263.50 ± 44.16	269.50 ± 42.78	7.53 ± 1.55 ª					

Results represent mean \pm SD. ^a Statistically significant result (p < 0.01).

Table 25 Serum proinflammatory cytokines and leptin concentrations in prepubertal obese children compared with controls (acc. to Aygun et $al^{[634]}$; with own modification).

Parameter	Obese children	Controls	P value
Leptin (ng/mL)	19.9 ± 7.4	7.9 ± 5.1	< 0.001
IL-1β (pg/mL)	33 ± 8.9	3.6 ± 1	< 0.001
IL-2 (U/L)	0.4 ± 0.1	0.9 ± 0.1	< 0.01
IL-6 (pg/mL)	45.2 ± 11.8	13.1 ± 3.9	< 0.001
TNF-α (pg/mL)	9.2 ± 2.3	3.9 ± 1	< 0.001
E-selectine (ng/mL)	78 ± 38	59 ± 29	< 0.01
hsCRP (mg/L)	4.1 ± 4.8	0.9 ± 1.5	< 0.001

Results are mean \pm SD; CRP, C-reactive protein; hs, high-sensitivity.

Monocyte/macrophages express CatL implicated in atherogenesis and associated arterial remodeling. Lipoprotein modification and uptake by atherosclerotic lesion cells, mainly macrophages and VSMCs, are important pathological steps in the formation of atherosclerotic plaques^[669,673,675,677,680-683]. CatL^[680,684,685] and CatB^[686,687] expression is enhanced in human coronary atherosclerotic lesions, carotid lesions, and in abdominal aortic aneurysms^[676]. CatB, CatL and CatD, were found in macrophage-derived foam cells in lipid-rich plaque areas^[688,689]. Macrophages, VSMCs, and endothelial cells can mobilize cathepsins also extracellularly, for example, the CatB, L, S, and K participated in plaque extracellular proteolysis^[690].

The role of cysteine cathepsin proteases in lipid uptake, storage, and efflux has been partly elucidated^[675,677]. Cathepsins have the capability to degrade LDL and reduce cholesterol efflux from macrophages, aggravating foam cell formation^[677]. Several cathepsins have been implicated in apoB-100 proteolytic modification, which enhances extracellular LDL particle aggregation, lipid droplet formation, and LDL retention to arterial proteoglycans^[683]. After taking up the lipoproteins and modified lipoproteins, macrophages and VSMCs become foam cells filled with lipid droplets. These lipids or modified lipoproteins affect cysteine cathepsin cellular expression and localization. Oxidized lipids translocate CatL and CatB to the cytosol and nuclei, which is consistent with the findings that lysosomal CatL and CatB localize to the cytoplasm and nuclei of apoptotic macrophages/foam cells in human carotid atheroma^[669,691].

Macrophages exposed to oxLDL or 7β -hydroxycholesterol, expressed high levels of CatB and CatL because of lysosomal destabilization, in addition to forming foam cells^[669]. The cathepsins translocated from lysosomes to cytosol or nuclei, caused foam cell apoptosis in the development and progression of atheroma^[673]. In addition, lysosomal cell permeabilization due to exogenous reactive oxygen species may induce lysosomal leakage leading to release of cathepsins into the cytoplasm^[675,692]. Cysteine cathepsins participated in the degradation of HDLs, thus reducing macrophage foam cell cholesterol efflux^[675].

CatB has been implicated in the pathogenesis of inflammatory diseases and in many aspects of atherosclerosis [677,693]. CatB, CatC, and CatL showed increased activity in the human aneurysm wall and thrombus [694-696], and CatB mRNA levels were increased in unstable regions of human plaque [687]. CatB cathepsin has been shown to activate IL-1 β converting enzyme caspase-1, important for secretion of the mature IL-1 β implicated in the pathogenesis of atherosclerosis [663,697].

CatC was involved in immunological response because it activated granzymes A and B, neutrophil elastase, and $CatG^{[698]}$, and was found to be upregulated in $CD8^+T$ cells in the thymus and spleen $^{[699]}$. Interestingly, CatC can be activated by CatL or CatS, but not by autocatalytic processing $^{[700]}$.

CatL is one of the most potent collagenases and elastases cleaving mature insoluble elastin [678]. This proatherogenic cathepsin is involved in death of macrophages, necrotic core formation and development of atherosclerotic plaque instability [701]. CatL is expressed in macrophages, VSMCs, and endothelial cells, and is located in the fibrous cap, tunica media, and macrophage-rich shoulder regions [676]. Macrophage apoptosis was found to be significantly correlated with expression of CatL in cell nuclei and membranes, and it was established that this cathepsin and TGF- β were overexpressed in cells subjected to mitochondrial metabolic stress [702].

CatL stimulated autophagy and inhibited apoptosis of oxLDL-induced endothelial cells^[703]. Thus, the proatherogenic effect of CatL was partly neutralized by inducing autophagy and inhibiting

apoptosis in early stages of atherosclerosis^[703]. Wei *et al*^[703] showed that oxLDL increased CatL protein expression and activation, inducing endothelial cell autophagy and apoptosis, and increasing endothelial cell permeability. The enhanced permeability of the endothelium is the first event in the cascade of bioprocesses leading to atherosclerotic formation, which leads to lipid infiltration and accumulation within the arterial wall^[703]. Inhibiting apoptosis lowered the permeability of monolayers to LDL^[704].

All together, cysteine protease cathepsins, such as for example CatB, CatC, and CatL, have been shown to play a critical role in immunity and development of lesions in atherosclerosis.

T. gondii infection

Cysteine proteinases play a major role in invasion and intracellular survival of a number of pathogenic parasites, including T. $gondii^{705}$. Cathepsin proteases act clasically as lysosomal hydrolases that digest endogenous and exogenous endocytosed polypeptides^[706]. They function in microneme and rhoptry protein maturation, host cell invasion, replication, and nutrient acquisition^[707,708]. Que $et\ al^{[705]}$ showed that the TgCatB, toxopain-1, was localized to rhoptries (secretory organelles required for parasite invasion into cells), and was critical for parasite invasion and rhoptry protein processing. Toxopain-1 was also found to be critical for infection in a chicken model of congenital toxoplasmosis^[709].

The parasite has a limited number of cystein proteinases, with one cathepsin B (TgCPB), one cathepsin L (TgCPL) and three cathepsins C (TgCPC1, 2, 3)^[710]. It was demonstrated that T: gondii tachyzoites ingested host cytosolic proteins and digested them using CatL and other proteases within its endolysosomal system^[708]. Both virulent type I and avirulent type II strain parasites ingested and digested host-derived proteins, indicating that the pathway is not restricted only to highly virulent strains. Larson $et\ al^{[711]}$ established that TgCPL was found in multiple puncts throughout the cytoplasm of intracellular replicating parasites, and was associated with a discrete vesicular structure in the apical region of extracellular parasites.

Cathepsins are critical to the parasite growth and differentiation^[707]. *T. gondii* cathepsins were required for peptide degradation in the parasitophorous vacuole (PV), as the degradation of the marker protein, *E. coli* β-lactamase, secreted into the PV of transgenic tachyzoites was completely inhibited by the CatC inhibitor^[707]. **Tables 26 and 27** summarized classes and specific cathepsin-like proteases of *T. gondii* proteases, their function and localization, and the parasite stage expressions^[708,712]. *Tg*CPC1 was found to be the most highly expressed cathepsin mRNA in tachyzoites, and three cathepsins, *Tg*CPB, *Tg*CPC1 and *Tg*CPC2, were undetectable in *in vivo* bradyzoites^[708].

Infection of the host vascular endothelial cells by *T. gondii* tachyzoites may be an important source of additional cathepsins activity, which will be superimposing on the host cell normal/pathologic cathepsin activities. It should be also added that, for example, CatB suppresses proliferation of peripheral blood mononuclear cells^[718], and therefore participates in the immunosuppressive activity of the parasite proteases that can lead to impairment of the host immune defense. As a result, *T. gondii* infection may play a crucial role in triggering, maintaining and progression of atherosclerosis, especially because of the rapid proliferation of tachyzoites in endothelial cells^[43], and global burden of congenital and acquired toxoplasmosis^[25,30].

Finally, it must be noted that there was a prevalence and pathology of amyloid in atherosclerotic arteries^[719], and active forms of Cat B, CatL, and CatD, were found to be associated with amyloid deposits

Table 26 Proteases characterized in T. gondii, their classes, function and localization (acc. to Li et $all^{7/2}$; with own modification).

Protease class	Specific protease	Protease function and localization		
	TgCPB/Toxopain-1	Invasion; rhoptry protein processing		
Cysteine	TgCPCs	Growth and replication; cleavage of dipeptides from N-terminal in cytosol	[707]	
TgCPL		Invasion; maturation of micronemal proteins; localized to the vacuolar compartment		
Metalloproteinase	Toxolysin 4	Invasion/egress; localized to the micronemes		
	TgSUB1	Invasion; maturation of rhoptry proteins; host cell attachment; localized to the micronemes	[715]	
Serine	TgSUB2	Invasion; maturation of rhoptry proteins; localized to the rhoptries	[716]	
	TgROM2,4,5	Surface protein sheddase; localized to tachyzoite plasma membrane	[717]	

Table 27 Features of *T. gondii* cathepsin-like proteases (acc. to Dou and Carruthers^[708]; with own modification).

currents , with own mounted only.				
Cathepsin protease	Mature molecular weight (kDa)	Activity	T. gondii stage expression	
TgCPL	30	endopeptidase	tachyzoite/bradyzoite	
ТдСРВ	28	endo/ exopeptidase	tachyzoite/bradyzoite	
TgCPC1	35	exopeptidase	tachyzoite	
TgCPC2	44	exopeptidase	tachyzoite	
TgCPC3	32	exopeptidase	sporozoite	

TgCPC1 was found to be the most highly expressed cathepsin mRNA in tachyzoites, and three cathepsins, TgCPB, TgCPC1 and TgCPC2, were undetectable in in vivo bradyzoites^[708].

in Alzheimer's disease^[720-722]. These findings are compatible with our earlier suggestions that latent chronic *T. gondii* infection may be at least in part responsible for the generation of amyloid plaques characteristic for the brains of patients with Alzheimer's disease and Down's syndrome^[222,223].

In summary, infection of host vascular endothelial cells by *T. gondii* tachyzoites may become an additional source of cysteine cathepsin activities that are superimposing on the activity of similar enzymes physiopathologically present in the host cells, because the parasite's tachyzoites, bradyzoites, and sporozoites express several members of these proteases.

TREATMENT WITH GLUCOCORTICOIDS (GCs) ACCELERATED AND/OR EXACER-BATED DEVELOPMENT OF ATHEROSCLE-ROSIS AT LEAST IN PART BY ENHANCING PROLIFERATION OF T. gondii TACHYZOITES IN MACROPHAGES AND FOAM CELL FORMATION

Atherosclerosis

Prolonged chronic stress associated with infection and inflammation can upregulate the levels of GCs and catecholamines leading to various diseases^[723], and specifically cardiovascular diseases have been associated with elevated levels of IL-1β, TNF-α, IL-6, and IFN-γ^[724]. GCs have been reported to interfere with a variety of both intra- and extracellular activities of mononuclear phagocytes, and GC receptors have been identified in monocytes and macrophages^[725]. Hydrocortisone and prednisone diminished human monocyte chemotaxis^[726], bactericidal capacity^[727,728], receptor activity for IgG and C3 complement^[727], and specifically inhibited macrophage elastase, collagenase, and PAI secretion^[729]. Moreover, GCs inhibited the expression and action of most cytokines^[730]. These drugs exerted an antiinflammatory effect by down regulating the production and secretion of proinflammatory cytokines such as TNF-α, IL-1β, IL-1β, IL-1β, IL-1β, II-1β.

2, IL-6, and the chemokines macrophage inhibitory protein (MIP)-1, and macrophage chemotactic protein/s (MCP)-1 and 2 from epithelial cells and macrophages, dendritic cells, T cells, and through apoptosis of T cells^[731,732]. GCs not only down-regulated proinflammatory cytokines, they also generated a functionally active, antiinflammatory phenotype in human monocytes that suppressed inflammatory processes and, thus, induced resolution of inflammation^[733].

Many studies provided the experimental and clinical evidence that treatment with GCs accelerated and/or exacerbated development of atherosclerosis^[194,734,735]. Accumulating evidence had demonstrated that foam cells were formed from macrophages *in vitro* at high concentrations (80-160 μg/mL) of oxLDL^[736,737]. Moreover, several authors showed that dexamethasone (DXM, 1μM) (a synthetic GC receptor agonist) could promote the macrophage-derived foam cell formation (a hallmark of early atherosclerotic lesions^[738,739]) at lower concentration of ox-LDL *in vitro*, and that the drug increased the formation of CEs in macrophages and human SMCs in a dose-dependent manner^[201,740]. The finding that DXM could promote the foam cell formation at lower ox-LDL concentration might be partially caused by the up-regulation of ACAT1 gene expression at the transcriptional level and the enhancement of cholesterol esterification^[194].

T. gondii infection

Wang et al^[741] showed that the growth rate of T. gondii RH strain tachyzoites was significantly increased in the peritoneal macrophages of rats treated with GCs in vivo compared with control cells (Table 28). They also demonstrated a significant inhibition of NO production in the macrophages collected from the rats receiving treatment with GCs compared with the control animals (Table 29). In addition, treatment with GCs inhibited the expression of iNOS mRNA in rat peritoneal macrophages^[741]. [Nb. Costa et al^[742] showed that T. gondii infection slightly upregulated glucocorticoid-induced TNF receptor expression in Treg cells and B cells, but the most robust increment in expression was observed in macrophages and dendritic cells] These findings are consistent with the results obtained by Nishikawa et al^[743] that T. gondii decreased NO production in the peritoneal macrophages, and with the suggestion that the parasite could partially decrease NO production in the infected host cells and therefore escape the immune defense reaction in the host^[744].

Jones *et al*^[745] also found that DXM (the GC receptor ligand) significantly reduced LPS-induced macrophages production of NO. It was reported that DXM can inhibit iNOS expression in LPS-treated murine macrophages by destabilizing the mRNA transcript^[746]. The GC receptor has also been demonstrated to interact with NF-κB, which has a role in induction of iNOS, therefore providing another possible means of inhibiting NO production^[747]. In addition, LPS-induced IL-12 generation also could be downregulated by the GC. Masur *et al*^[748] observed that even therapeutic concentrations of hydrocortisone rendered normal macrophages unresponsive to IFN-γ

and prevented the induction of the activated mononuclear phagocyte the capacity to respond to *T. gondii* ingestion with an enhanced oxidative respiratory burst, and the ability to inhibit intracellular parasite replication.

GCs are widely used for the treatment of autoimmune diseases and after bone marrow transplantation, sometimes resulting in development of acute toxoplasmosis in these patients^[741,749,750]. Approximately one-third of transplant recipients experienced neuropsychiatric and other complications including encephalopathy, stroke, seizures, loss of vision, cardiovascular events, and de novo lymphoproliferative disorders (lymphoma, glioma)[751-754]. Organ transplant recipients had a three- to fourfold higher incidence of malignant diseases development compared with the general population^[752], and many of CNS lymphomas were associated with prior Ebstein Barr virus infections^[755]. Neurologic complications of liver transplantation were more common than that of other solid organ transplants (13-47%)^[753]. Recurrent ocular toxoplasmosis has also been reported in patients receiving systemic corticosteroid therapy (dose range, 0.27-1.23 mg/kg/day)[756]. The study of Ahlbom et al[757] suggested that prenatal exposure to excess of GCs increased the susceptibility of cerebellar granule cells to oxidative stress-induced cell death, and elevated maternal cortisol levels during pregnancy have been associated with reduced childhood $IQ^{[758]}$. It must be emphasized that in mice brains infected with T. gondii electron microscopy revealed that cortisone increased the amount of tachyzoites, cysts and cystozoites, as the rupture of cysts released a highly resistant antigen of cystozoite type^[759]. All these CNS abnormalities, clinical complications, and developmental disturbances observed during therapeutic use of GCs could be due to the increased proliferation of T. gondii in the host cells induced by the increased levels of GCs and other immunosuppressive drugs that markedly affected physiologic balance between profusely generated type $T_{\rm H}1$ and type $T_{\rm H}2$ cytokines $^{[760,\ 761]}.$

Taken together, several authors showed that DXM could promote the macrophage-derived foam cell formation at lower concentration of ox-LDL *in vitro*, and that the drug increased the formation of CEs in macrophages and human SMCs in a dose-dependent manner. These changes might be partially caused by the up-regulation of ACAT1 gene expression and thus the enhancement of cholesterol esterification. However, it should be noted that the growth rate of RH strain *T. gondii* tachyzoites was significantly increased in the peritoneal macrophages of rats treated with GCs *in vivo* compared with control cells. A significant inhibition of NO production in the macrophages collected from the rats receiving treatment with GCs was also observed. Moreover, GCs inhibited the expression and action of most cytokines involved in inflammation. Thus, all these molecular processes may at least in part explain an important role of GCs in development of atherosclerosis.

BENEFICIAL EFFECTS OF VITAMIN D IN BOTH ATHEROSCLEROSIS AND T. gondii INFECTION. VITAMIN D MARKEDLY DECREASED PROLIFERATION OF T. gondii TACHYZOITES IN MACROPHAGES AND REDUCED TISSUE PATHOLOGY IN ANIMALS INFECTED WITH THE PARASITE

Atherosclerosis

Vitamin D regulates wide range of physiological and pathological

Table 28 Proliferation of *T. gondii* tachyzoites in peritoneal macrophages of rats treated with glucocorticoids (GCs) (acc. to Wang *et al*^[21]; with own modification).

GCs	Number of T. gondii per 100 cells				
	1 hr	12 hrs	24 hrs		
DXM	35 ± 3	55 ± 8 ª	176 ± 15 ª		
HSS	39 ± 5	81 ± 12 ª	227 ± 14 ^a		
MP	37 ± 6	85 ± 14 ^a	242 ± 16 a		
Control	39 ± 5	23 ± 4	16 ± 4		

Results are expressed as means \pm SD, number of rats =3. $^{\rm a}$ p < 0.01 vs control. Rats were injected intramuscularly with a dose of GCs for 7 days, macrophages were harvested and cultured for 12 hrs and then incubated with T. gondii at the ratio 1:1 (parasites/macrophages = 1:1). DXM, dexamethasone; HSS, hydrocortisone sodium succinate; MP, methylprednisone.

Table 29 Nitrite production by peritoneal macrophages from rats treated with glucocorticoids (GCs) (acc. to Wang *et al*^[21]; with own modification).

GCs	Nitrite production (µM)				
	12 hrs	24 hrs	36 hrs		
DXM	9.78 ± 0.37 a	14.31 ± 1.22 a	17.99 ± 1.08 a		
HSS	8.19 ± 0.32 a	11.05 ± 0.74 a	13.51 ± 1.45 a		
MP	7.86 ± 0.46 a	10.65 ± 0.98 a	12.90 ± 0.99 a		
Control	14.44 ± 1.04	25.41 ± 1.44	30.85 ± 1.62		

Results are expressed as means \pm SD, number of rats = 3. $^{\rm a} p < 0.01~vs$. control. DXM, dexamethasone; HSS, hydrocortisone sodium succinate; MP, methylprednisone. Peritoneal macrophages isolated from rats were treated with GCs for 7 days, and then incubated for 12, 24, and 36 hrs. NO₂ production in the supernatant of the cell culture medium was measured by the Griess reaction.

processes characteristic for atherosclerosis, including vascular cell growth, migration, and differentiation, immune response modulation, cytokine expression, and inflammatory and fibrotic pathways[762]. Vitamin D deficiency is affecting more than one billion people to approximately 50% of population worldwide^[762-764]. Shortage of vitamin D induced atherosclerosis and high blood pressure in mice^[765]. Epidemiological and clinical evidence linked vitamin D deficiency to cardiovascular diseases in humans^[766], including carotid atherosclerosis[762,767,768]. Vitamin D receptor (VDR) is present in all cells implicated with atherosclerosis, such as endothelial cells, vascular smooth muscle cells, and immune cells[762]. It was demonstrated that vitamin D inhibited foam cell formation by suppressing oxidized and acetylated LDL-derived cholesterol uptake in macrophages from type 2 diabetes patients^[769], and decreased atherosclerosis by regulating T lymphocyte functions^[770]. Vitamin D exerted also protective effects on endothelial cell dysfunction, inflammatory processes that precede atherosclerosis^[771], regulated the expression of profibrotic and antifibrotic factors[772], and inhibited formation of atherosclerosis calcification^[773,774].

Vitamin D was found to modulate both the innate and acquired immune systems^[775-778]. Macrophages, endothelial cells, and smooth muscle cells, among others, are able to transform 25(OH)₃D₃ to its active hormonal form, 1,25-dihydroxyvitamin D (1,25(OH)₂D₃). The active metabolite primarily mediated its effects through the intracellular VDR^[779], which is expressed in most cell types of the immune system, in particular in antigen presenting cells such as monocytes, macrophages, and in dendritic cells^[780], B cells, as well as in CD4⁺ and CD8⁺T cells^[775,778]. Vitamin D favors a mononuclear cell phenotype, increasing VDR expression on monocytes and macrophages^[779]. Activation of VDR altered transcription, proliferation, and differentiation of immune cells^[779], and modulated immune responses both indirectly, by reducing the activation of proinflammatory T cells by antigen-presenting cells^[781],

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and directly, by inhibiting T and B cells proliferation [775,782]. Active vitamin D and corticosteroids exerted additive immunosuppressive effects on $T_{\rm H}1$ responses [783]. All these actions finally result in a $T_{\rm H}2$ type-driven antiinflammatory state in the host [784,785]. **Table 30** listed multiple actions of vitamin D in the immune system. Several of the above-presented actions of vitamin D on immune system may have beneficial effects on clinical course and outcome of patients with atherosclerosis. It should be added that also vitamin A exerted favorable effects in patients with atherosclerosis because it decreased cytotoxicity of oxidized LDL and improved peripheral blood mononuclear cells viability [801].

T. gondii infection

An infectious pathogenesis put forward that intracellular parasite infection disrupt the vitamin D regulated immune system resulting in persistent infection and chronic inflammation. Studies showed that 1,25(OH)₂D₃ directly increased antimicrobial peptide gene expression which could enhance host defense against infection^[802,803]. Vitamin D was also required for IFN-γ-mediated antimicrobial activity of human macrophages^[804].

Vitamin D facilitate neutrophil motility and fagocytic function^[805]. Circulating vitamin D levels have a direct influence on macrophages, increase their "oxidative burst" potential, such as for example maturation and generation of cytokines, hydrogen peroxide, and acid phosphatase^[787,806]. Vitamin 1,25-D₃ inhibited proliferation of T_H1 cells through impairing production of IL-2, TNF-α, and IFN, as well as T_H17 cells, and thus skewing cytokine production toward a T_H2 phenotype^[787,807]. Vitamin D may thus improve outcomes by reducing both local and systemic inflammatory actions as a result of modulating cytokine responses and decreasing TLR activation^[808].

Finally, it must emphasized that vitamin D significantly decreased *in vitro* proliferation of *T. gondii* tachyzoites in macrophages (**Table 31**), increased NO generation in these cells (**Table 32**) [809], and diminished various tissue pathology in animals infected with the parasite (**Table 33**), possibly by acting on tachyzoites in parasitophorous vacuole [810]. Of note, vitamin D may however also be linked to the increased susceptibility and mortality of mice infected with the pathogen, probably because of its downregulation of the T_H1 type cytokine response [811].

In summary, experimental and clinical data provide vast evidence

that vitamin D have important beneficial impact on immunity, as well as on *T. gondii* infection, and therefore it should find a solid place in various treatment regimens directed against both atherosclerosis and the pathogen.

CONCLUDING REMARKS

Intracellular parasite T. gondii is widely disseminated in animals and infects approximately 30-50% of human population worldwide. The pathogen attacks almost all nucleated cells, and vascular endothelial cells have increased susceptibility to infection with T. gondii causing oxidative stress and endothelial dysfunction. Persistent inflammation associated with the increased production of proinflammatory cytokines and cysteine cathepsins during latent chronic toxoplasmosis cause enhanced generation of foam cells, disturbances of blood coagulation, and development of atherosclerotic lesions. In these bioprocesses, T. gondii tachyzoites become an additional source of TgACAT1 and TgACAT2 and cathepsin proteases that are superimposing on the respective enzymes produced in the host vascular endothelial cells, thereby enhancing their final activities. These findings may be supported by at least six facts: (a) early stages of atherosclerosis began already in early embryonic life probably caused by congenital toxoplasmosis acquired in the first trimester of gestation as a result of parasite transmission from mother to fetus[110]; (b) the parasite penetrated nucleated erythroblasts and macroreticulocytes from fetal mouse liver and the circulating erythrocytes of fetal mice[414], and multiplied in the immature cells[812]; (c) the accelerated atherosclerotic development was due to the increased levels of cholesterol and LDL reported in T. gondii infected humans^[16] and animals^[813]; (d) the highest amongst analyzed regression coefficients (B = 12.49, p = 0.026, Eta² = 0.058) between the prevalence of toxoplasmosis and cardiovascular (cerebrovascular and ischemic heart) diseases in a set of 88 countries[30]; (e) there is a frequent comorbidity of severe carotid atherosclerosis and mood disorders $(p < 0.0001)^{[814]}$ as well as a significant relationship between chronic T. gondii infection and mood disturbances[815]; and (f) pravastatin, simvastatin and other statins (drugs with hypocholesterolemic and antiatherosclerotic activities) inhibited the adhesion, replication and proliferation of T. gondii^[816-819]; propranolol, a β-adrenoreceptor blocking agent, also exerted antitoxoplasmic

Table 30 Effects of 1,25-dihydroxyvitamin D_3 on innate immune system (acc. to Pelajo et $all^{[786]}$; Youssef et $all^{[787]}$; with own modification).

Vitamin D actions in the immune system	Refs
Decreases the antigen-presenting activity of macrophages to lymphocytes; acts on phenotype and function of antigen presenting cells (monocytes, macrophages, dendritic cells) via oxidative burst, production of acid phosphatase and hydrogen peroxide, and maturation of cytokines; acts on neutrophils mobility and phagocytic function; upregulates antimicrobial peptides (cathelicidins, beta-2/beta-3 defensins) in neutrophils, NK cells, monocytes, macrophages, and lymphocytes	[787-789]
Inhibits the maturation of monocytes into dendritic cells	[790, 791]
Induces the activation of T reg and NK T cells	[792, 793]
Inhibits TH1 type cytokine response	[783, 790, 791, 793]
Decreases IL-2, IFN-γ, and CD4 and CD8 DNA synthesis	[776, 787-791]
Stimulates the T _H 2 type cytokine dominance	[783,790, 791, 793]
Increases IL-4, IL-5 and IL-10 synthesis	[777, 789, 790]
Inhibits the synthesis of IL-12, IL-1, IL-6 and TNF- α	[776, 788, 794, 795]
Inhibits B cell proliferation, plasma cell differentiation, and antibody/immunoglobulin production	[789-791, 796, 797]
Triggers B cell apoptosis	[787]
Increases apoptosis induced by dendritic cells and T lymphocytes – tolerance	[797]
Induces autophagy in human monocytes/macrophages via cathelicidin	[777, 780]
Induces NOS in a human monocyte/macrophage cell line ^a	[798]

NOS, nitric oxide synthase. *NO is an effector molecule of parasite killing^[799]. Interestingly, $1\alpha25$ (OH)₂D₃ suppressed growth and triggered the destruction and clearance of another intracellular pathogen *Mycobacterium tuberculosis* by activating autophagy in infected human monocytes/macrophages^[800].

Table 31 Effect of vitamin D_3 and IFN- γ on proliferation of *T. gondii* (RH strain) tachyzoites per infected peritoneal macrophage of BALB/c mice after incubation for 96 hrs in RPMI1640 cells culture (acc. to Ghaffarifar *et al.*^[809]; with own modification).

Experiment No.	Controls	Solvent ^a	Vit D ₃ (1000 IU)	IFN-γ (100 IU)	Vit D ₃ (1000 IU) plus IFN-γ (100 IU)
1	3.01 ± 0.14	2.93 ± 0.16	2.49 ± 0.19 ^b	2.6 ± 0.2 ^b	2.37 ± 0.19 ^b
2	3.15 ± 0.12	3.03 ± 0.16	2.74 ± 0.16	2.5 ± 0.15 ^b	2.58 ± 0.13 ^b
3	3.05 ± 0.15	3.04 ± 0.14	2.82 ± 0.17	2.57 ± 0.16 ^b	2.69 ± 0.2 ^b
4	3.16 ± 0.14	3.0 ± 0.14	2.39 ± 0.19 ^b	2.59 ± 0.2 ^b	2.03 ± 0.19 ^b

Numbers of tachyzoites are given as a mean \pm SD. ^a Ethanol 95. ^b Statistically significant differences compared with controls ($p \le 0.05$).

Table 32 Effect of vitamin D_3 and IFN- γ on NO production by peritoneal macrophages of BALB/c mice infected with *T. gondii* (RH strain) after incubation for 24 hrs in RPMI1640 cells culture (acc. to Ghaffarifar *et al*^[809]; with own modification).

Experiment No.	Controls	Solvent ^a	Vit D ₃ (1000 IU)	IFN-γ (100 IU)	Vit D ₃ (1000 IU) plus IFN-γ (100 IU)
1	109 ± 8.02	108.2 ± 12.45	165 ± 11.30 ^b	146 ± 7.22 ^b	187.8 ± 9.82 ^ь
2	108 ± 9.46	108.9 ± 6.93	121.2 ± 6.68	139.5 ± 5.76 ^b	136.2 ± 10.21 ^b
3	109.6 ± 7.35	108.2 ± 4.96	139 ± 7.01 ^b	146 ± 4.93 ^b	146.9 ± 9.62 ^b
4	109 ± 7.03	108.6 ± 4.26	166 ± 7.01 ^b	146.2 ± 5.60 ^b	191.5 ± 9.62 ^b

Values are given as mean \pm SD. ^aEthanol 95. ^bStatistically significant results compared with controls ($p \le 0.05$). NO production was estimated as a nitrite release from infected macrophages (μ M).

Table 33 Effect of pretreatment with $1,25(OH)_2D_3$ (0.5 $\mu g/kg/2$ days) on tissue pathology caused by *T. gondii* avirulent ME49 strain infection with 20 cysts administered intraperitoneally in BALB/c mice (acc. to Rajapakse *et al*^[810]; with own modification).

Tissue	Pathology	No treatment	Treatment with Vit D ₃
Luna	Alveolar macrophages	1	0
Lung	Inflammatory foci	2	1
	Inflammatory foci	3	2
Liver	Hemorrhage	2	0
	Mitosis	1	0
Small intestine	Inflammatory infiltrates	1	0
Small intestine	Necrotic mucosal cells	2	1
Brain	Presence of the parasite	2	0
Spleen	Granulocytes	2	1

Histopathologic examination of the tissues was performed 7 days post inoculation. Numbers are based on severity of the lesions (0, no lesion, 1, mild, 2, slight, 3, moderate changes) and the total was divided the number of animals in the group. Also, in vitro studies with incubated intestinal epithelial cells showed a significant dose-dependent inhibition of intracellular *T. gondii* tachyzoites (RH strain, type I) proliferation at 10⁻⁷ M of 1,25(OH)₂D₃ concentration.

effects^[820,821] and had both antiatherosclerotic^[822] and atherogenic^[823] properties. Thus, it seems that latent chronic toxoplasmosis play an essential but so far neglected role in the pathogenesis of atherosclerosis.

Finally, one may ask why until now there was no suggestion that this pathogen might be so crucial and successful environmental factor responsible for triggering and development of atherosclerosis on a global scale? Probably, the answer to this question refers to the facts that physicians have a very limited knowledge about this widely disseminated pathogen in humans, animals and birds, and there is a general opinion that such infection is usually harmless in immunocompetent individuals^[28,29]. Moreover, there is no sufficiently specific and sensitive laboratory method available so far for diagnosis of infection with the parasite, and a plethora of T. gondii antigens detected in human plasma due to various parasite strains that infect host, and different stage-specific antigens of the pathogen^[824]. T. gondii antigens recognized by IgG antibodies also differ between immunocompetent hosts with and without active proliferation of tachyzoites during chronic infection[825]. Recently however, we proposed and gave the reasons^[495] for a need to use a very specific method proposed long time ago by Sternberger et al^[826] and modified by Conley and Jenkins[827].

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