Paraoxonase (PON)-1: a brief overview on genetics, structure, polymorphisms and clinical relevance

Nelusha Shunmoogam¹
Poobalan Naidoo¹
Robert Chilton²

¹Sanofi, Midrand, South Africa; ²Department of Medicine, Division of Cardiology and Interventional Cardiology, University of Texas Health Science Center at San Antonio, San Antonio, TX, USA

Abstract: Paraoxonase-1 (PON1) is a high-density lipoprotein-associated esterase and is speculated to play a role in several human diseases including diabetes mellitus and atherosclerosis. Low PON1 activity has been associated with increased risk of major cardiovascular events, therefore a variety of studies have been conducted to establish the cardioprotective properties and clinical relevance of PON1. The major aim of this review was to highlight the important studies and to subsequently assess if PON1 has clinical relevance. A review of the literature showed that there is currently insufficient data to suggest that PON1 has clinical relevance. It is our opinion that robust studies are required to clarify the clinical relevance of PON1.

Keywords: antioxidant, inflammation, atherosclerosis, cardiovascular disease

Introduction

Human serum paraoxonase-1 (PON1) is a calcium-dependent hydrolytic enzyme that is found in a variety of mammalian species. Abraham Mazur¹ and Norman Aldridge² played a pivotal role in the identification and classification of PON1 in the mid-1940s to early 1950s. Initially, the enzymes were referred to as “A”-esteras, but later became universally known as paraoxonases due to their ability to detoxify the organophosphate compound paraaxon which is the toxic metabolite of parathion, a commonly used agricultural insecticide.³

PON1 belongs to a family of three serum paraoxonases, including PON2 and PON3; however, PON1 remains the most popular member of this family.⁴ This is largely due to the elegant studies by Mackness et al that described the role of high-density lipoprotein (HDL)-associated PON1 in decreasing lipid peroxide accumulation on low-density lipoprotein (LDL).⁵⁶⁻⁷ This highlighted a link to PON1 and cardiovascular disease, which sparked the research interest in PON1, mainly to elucidate the precise physiological mechanisms of the enzyme. In addition, PON1 hydrolyzes homocysteine thiolactone. Homocysteine thiolactonase activity of PON1 protects against N-homocysteinylation, which is detrimental to protein structure and function.⁸

Structure of PON1

PON1 is a calcium-dependent enzyme consisting of 354 amino acids with a molecular mass of 43 kDa. Structural analysis using X-ray crystallography revealed the six-bladed β-propeller structure of PON1 (Figure 1), with a central tunnel that houses two calcium ions.⁹ Each calcium ion, depending on its location within the enzyme, plays an important part in the activity of PON1.¹⁰ The calcium ion located deeper within
found associated with HDL in the plasma.12,13 PON1 forms and PON3 primarily synthesized in the liver and mostly are expressed in various mammalian tissues, with PON1 PON1–HDL interactions.9

Helix 3. Reproduced from Blaha-Nelson

H1, H2 and H3, where H1 and H2 have functions in activation. Three helices are located above the active site of and is important for substrate positioning and ester bond lies at the bottom of the active site cavity has a catalytic role in PON1–HDL interactions.9 The other calcium ion which stabilizes PON1. The other calcium ion which

the tunnel has a structural role that is critical for the conformational stability of PON1. The other calcium ion which lies at the bottom of the active site cavity has a catalytic role and is important for substrate positioning and ester bond activation. Three helices are located above the active site of PON1: H1, H2 and H3, where H1 and H2 have functions in PON1–HDL interactions.9

Genetics of PON1

The human PON1 gene is a member of a multigene family consisting of three members in total. PON1, PON2 and PON3 are located next to each other on chromosome 7 and share extensive structural homology. Interestingly, PON1 can be differentiated from PON2 and PON3 by the three extra nucleotide residues in exon 4.11 The genes for this family are expressed in various mammalian tissues, with PON1 and PON3 primarily synthesized in the liver and mostly found associated with HDL in the plasma.12,13 PON1 forms part of a repertoire of HDL-associated enzymes, including lecithin-cholesterol acyltransferase and platelet-activating factor acetyl-hydrolase, responsible for the antioxidative activity of HDL.

PON1 polymorphisms

Human PON1 has many single-nucleotide polymorphisms (SNPs); eight have been identified on the promoter region and 176 within the gene sequence,14 some of which exert changes in PON1 level and activity. These polymorphisms may also affect the risk for disease development and the severity of disease.15 Studies have identified two common polymorphisms in the coding region (at positions 55 and 192) that have been reported to affect the activity and concentration of PON1.16

The leucine/methionine polymorphism at position 55 of the amino acid sequence (L55M) has been associated with changes in PON1 serum concentrations, and an association with the occurrence of cardiovascular disease was also observed.17 The glutamine/arginine polymorphism at position 192 (Q192R) has been shown to affect PON1 activity, where the Q192 isoform was demonstrated to hydrolyze paraoxon and metabolize oxidized LDL more effectively than the R192 isoform.18 The Q192R polymorphism is regarded as the chief biomarker of oxidant status, where LDL oxidation is prevented the most in QQ homozygous patients and the least in RR patients.19

In addition, three common SNPs (G-907C, A-162G and C-108T) were identified in the promoter sequence of the PON1 gene.14 These polymorphisms are associated with considerable differences in PON1 concentration and activity, and they have also been implicated in the presence of coronary heart disease.20 It has been well established that low PON1 activity is linked with an increased risk of cardiovascular disease, implicating PON1 as a physiologically important enzyme.

Clinical relevance

PON1 is an HDL-associated protein that has the ability to hydrolyze oxidized LDL-cholesterol, with potential atheroprotective effects.21 Furthermore, PON1 can cleave phospholipid peroxidation adducts with potential cytoprotective functions.22 Given these potential atheroprotective effects, and the large burden of atherosclerotic cardiovascular disease, considerable work has focused on elucidating the clinical relevance of PON1.

Animal studies23,24 have suggested atheroprotective benefits of PON1. Transgenic mice overproducing human PON1 protected them from atherosclerosis, when compared to wild-type mice.23 In addition, PON1-deficient mice are at greater risk of developing atherosclerosis than wild-type mice.24 Animal studies have various limitations and direct extrapolation to humans cannot be made. However, they are useful for “transiting” from in vitro studies to clinical studies.

With respect to clinical studies, Mackness et al25 investigated the effects of the C-108T and G-909C promoter polymorphisms on PON1 levels and the presence of coronary heart disease (CHD). It was a case–control study, with 417 people with CHD and 282 healthy controls. PON1 activity and concentration were significantly lower in the
CHD population compared to controls, regardless of their C-108T and G-909C genotype \( (p<0.001) \). Both promoter polymorphisms were not associated with CHD presence and the authors concluded that PON1 status was significantly lower in people with CHD.

Azarsiz et al\(^23\) investigated PON1 activity in patients with angiographically confirmed CHD. Twenty-four healthy volunteers and 101 patients were enrolled in the study; 68 patients had coronary artery disease, which was confirmed by coronary angiography. PON1 activities of patients with CHD were lower than those of controls; however, this difference was statistically insignificant. In contrast to Azarsiz et al’s study,\(^25\) Sharma et al\(^26\) demonstrated that serum PON1 activity was significantly low in CHD patients. Sharma et al investigated PON1 activity in 200 patients suffering from CHD and 150 normal individuals. CHD patients were classified into two groups on the basis of associated risk factors (diabetes mellitus, hypertension): group 1 \( (n=120; \) CHD patients with associated risk factors) and group 2 \( (n=80; \) CHD patients with no associated risk factors).\(^26\)

Smoking is an important independent risk factor for CHD, and a study by Han et al\(^27\) showed that cigarette smoking status together with the presence of common PON1 SNPs play a role in the risk of developing CHD.\(^27\) In this case-control study nested within the Singapore Chinese Health Study, which was a prospective cohort of 63,257 participants recruited over 5 years, Han et al evaluated 1,914 Singaporean Chinese: 688 cases and 1,226 controls. The participants were stratified according to cigarette smoking status as ever-smokers \( (n=813) \), who answered yes to smoking at least one cigarette a day for a year or longer, and never-smokers \( (n=1,101) \). The T allele of the PON1 rs6626 polymorphism was shown to be associated with an increased risk of CHD among ever-smokers only \( (\text{OR}=1.35, \ 95\% \ CI \ 1.08–1.68; \ \text{adjusted} \ p=0.036) \), whereas another PON1 SNP, rs3735590, was shown to be associated with an increase in CHD among never-smokers only \( (\text{OR}=1.53, \ 95\% \ CI \ 1.11–2.11; \ \text{adjusted} \ p=0.036) \),\(^27\) highlighting that more research is required on PON1 polymorphisms and CHD.

Kunutsor et al\(^29\) investigated the association of PON1 with cardiovascular risk. The study included prospectively measuring PON1 activity in 6,902 study subjects. The study subjects were followed up for a mean of 9.3 years, with 730 adverse cardiovascular events being recorded. There is an approximately log-linear inverse association between PON1 activity and CVD risk, which is partly dependent on HDL-cholesterol levels. Kunutsor et al did a further meta-analysis of six studies with 15,064 study subjects and 2,958 incident adverse cardiovascular events. Based on the findings of Kunutsor et al, although there is a negative relationship between PON1 and adverse cardiovascular events, PON1 does not add value to cardiovascular risk stratification, beyond conventional cardiovascular risk factors.\(^28\)

Given that atherosclerotic cardiovascular disease is a major cause of mortality, it was hypothesized that older individuals may have lower PON1 activity, when compared to younger individuals. Seres et al\(^29\) investigated the relationship between age and serum PON1 activity. One hundred twenty-nine healthy subjects aged between 22 and 89 years were included in their study. Serum PON1 activity significantly decreased with age \( (r=-0.38, \ p<0.0001) \). HDL concentrations remained unchanged with age; however, Apo A1 concentration showed a slight negative, but significant correlation with age \( (r=-0.19, \ p<0.027) \). Moreover, the total cholesterol concentration was positively and significantly correlated with age \( (r=0.40, \ p<0.001) \). The authors also noted that HDL from elderly subjects was more susceptible to oxidation than HDL from young subjects, measured by higher lipid peroxidation rate. The study was limited by relatively small sample size, but did demonstrate reduction in PON1 activity with age.

Numerous further studies investigated whether PON1 is a longevity gene. Lescai et al\(^30\) carried out a meta-analysis of these aforementioned studies that included 5,962 subjects: 2,795 young controls \( (<65 \text{ years of age}) \) and 3,167 old subjects \( (>65 \text{ years of age}) \). R carriers demonstrated a significant result with an overall OR of 1.16 \( (95\% \ CI \ 1.04–1.30, \ p=0.006) \). The QR genotype also showed a significant result, with an overall OR of 1.14 \( (95\% \ CI \ 1.02–1.27, \ p=0.016) \). The authors concluded that PON1 gene variants at codon 192 impact on the probability of attaining longevity; and those subjects carrying RR and QR genotypes \( (R+ \text{ carriers}) \) are favored in reaching extreme ages. However, subsequent meta-analysis with larger number of patients\(^31,32\) has suggested that there is no effect of PON1 on human longevity. However, population-specific effects could not be excluded. The aforementioned meta-analysis may be limited by publication bias and variations in analytical procedures used to measure PON1 activity.

Another meta-analysis based on 30 publications analyzed the risk of cancer in relation to the PON1 Q192R polymorphism.\(^33\) Considering PON1 polymorphisms have been associated with several types of cancer, Zhang et al investigated the role of the genetic variant Q192R in cancer susceptibility. The authors conducted a thorough search of the literature and based on robust inclusion criteria, their meta-analysis exam-
ined 30 publications with a total of 8,112 cases and 10,037 controls. The results indicated that the PON1-192 R allele was associated with a reduced risk of overall cancers compared to the 192 Q allele (OR=0.842, 95% CI 0.725–0.979); however, when the results were analyzed according to the cancer types, an increased and decreased risk of cancer subtypes were observed under heterozygous, homozygous, dominant and recessive models. It is well established that oxidative stress and increased free radicals may lead to an increased risk of cancer; therefore, the antioxidant properties of the genetic variants of PON1 should be studied in more detail to fully understand their role in cancer.

Diabetes mellitus is characterized by increased oxidative stress and damage, possibly due to the result of glycosylation of LDL by glucose. Various studies have demonstrated a reduction in PON1 in type 2 diabetic patients. Furthermore, reduced PON1 activity in type 2 diabetes mellitus has been associated with increased risk of cardiovascular disease. Rozek et al postulated that reduced PON1 activity in diabetic patients results in reduced HDL protective activity against cell membrane peroxidation contributing to increased arteriosclerosis in diabetic patients. Studies demonstrating reduction in PON1 levels in diabetic patients are contradicted by studies that show no changes in the levels of PON1 in diabetic subjects. However, although the aforementioned studies showed no absolute reductions in PON1 levels, they demonstrated qualitative reductions in PON1 activity.

Nie et al conducted a meta-analysis on the relationship of PON genes and Alzheimer’s disease. Fifteen studies (involving five polymorphisms) were included in the meta-analysis. The authors concluded that the “SS genotype of PON2 S311C polymorphism had a significant association with Alzheimer’s disease in the studied population, and the A allele of PON1 rs705379 polymorphism was positively related to AD in the Caucasian population as well as the GG genotype decreased AD risk significantly in Caucasians.” The meta-analysis is limited by the quality of studies included; further robust studies are required to elucidate the role of PON in Alzheimer’s disease.

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A brief overview of paraoxonase (PON)-1

Conclusion
Although the preclinical and clinical studies around PON1 are intriguing, there is currently insufficient data to suggest that PON1 has clinical relevance. Furthermore, the study findings are incongruent and, in some instances, contradictory. Robust studies are required to clarify the clinical relevance of PON1.

Disclosure
PN is employed by Sanofi and NS is a previous employee of Sanofi. Sanofi has molecules in the lipid, antiplatelet, hypertension, diabetes, anticancer, anti-infective and anti-depressant therapeutic areas. Neither PN nor NS own stocks in Sanofi. The other author reports no conflicts of interest in this work.

References


