Original article

Effects of acetylsalicylic acid alone or with omega-3 in patients with chronic coronary artery disease

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\textbf{A B S T R A C T}

Introduction and objectives: Antiplatelet agents such as acetylsalicylic acid (ASA) play a prominent role in preventing atherothrombosis. However, low-responsive patients who will not benefit from an increased dosage of this drug, which can cause bleeding and gastrointestinal irritation, need to be identified. Drugs such as omega-3 fatty acids, which enhance the vasodilating condition and diminish platelet aggregation, can potentiate the antiaggregating effects of ASA, avoiding its side effects. Thus, we assessed the alternative use of 200 mg/day of ASA and 100 mg/day of this drug combined with 1 g of omega-3 in 152 patients with chronic coronary artery disease.

Methods: Our analysis included platelet function (ASPItest), TBX2 concentrations (ELISA), and SNPs polymorphisms in the rs3842787 and rs3842798 regions of the PTGS1 gene of the COX-1 enzyme and the rs5918 region of the ITGB3 gene of the fibrinogen’s receptor subunit glycoprotein IIIa.

Results: ASPItest detected 38 non-responders. The reduction of ASPItest values was more significant in this group than in responders and fell to levels of responders in non-responders of the 200 mg/day treatment. A rare allele of rs3842787 is associated with a worse ASPItest response, and the rare allele of the rs5918 polymorphism with a worse response related to TBX2 concentration. Both treatments showed no statistically significant difference in hematuria or bleeding, constituting safe treatment alternatives, and omega-3 treatment reduced monocyte levels.

Conclusions: Our results underscore the usefulness of pharmacogenetics for personalized treatments, avoiding gastrointestinal effects and undesirable bleeding.

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\textbf{Abbreviations:} ASA, acetylsalicylic acid; SNPs, single-nucleotide polymorphisms; TBX2, thromboxane B2; GPIIIa, glycoprotein IIIa; NR, non-responders’; R, responders’.

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\url{https://doi.org/10.1016/j.rccl.2023.07.002}

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Efectos del ácido acetilsalicílico solo o combinado con omega-3 en pacientes con enfermedad arterial crónica

RESUMEN

Introducción y objetivos: El ácido acetilsalicílico (AAS) es ampliamente utilizado para prevenir la trombosis arterial. Sin embargo, deben identificarse pacientes refractarios para los cuales dosis altas de AAS son ineficaces y pueden causar irritación gastrointestinal y hemorragia. El fármaco omega-3 puede aumentar los efectos anticoagulantes del AAS y evitar sus efectos colaterales. Por ese motivo evaluamos el uso de 200 mg/día de AAS o 100 mg/día de AAS combinado con 1 g de omega-3 en 152 pacientes con enfermedad coronaria crónica.

Métodos: Se analizaron la función plaquetaria (ASPtest), las concentraciones de TBX2 (ELISA) y polimorfismos de SNP en las regiones rs3842787 y rs3842798 del gen PTGS1 de la enzima COX1 y la región rs5918 del gen ITGB3 de la subunidad glicoproteína IIIa del receptor de fibrinógeno.

Resultados: El ASPtest detectó 38 pacientes refractarios. La reducción de la función plaquetaria fue más significativa en este grupo que en los respondedores, cayendo a niveles de respondedores en los refractarios al tratamiento con 200 mg/día. Un alelo raro de rs3842787 está asociado a una peor respuesta al ASPtest, y un alelo raro del polimorfismo rs5918, a una peor respuesta en el análisis de la concentración de TBX2. Los tratamientos analizados no mostraron evidencia estadísticamente significativa de hematuria o de sangrado, y el tratamiento con omega-3 mostró reducción en los niveles de monocitos.

Conclusiones: La farmacogenética es una herramienta indispensable para una medicina personalizada con tratamientos que eviten efectos gastrointestinales y hemorragias, como los descritos en este estudio.

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The refractory effect in ASA non-responsive individuals is due to several factors, among which the single-nucleotide polymorphisms (SNPs) stand out.16 Thus, pharmacogenetics can effectively contribute to identifying and treating ASA-responsive and low-responsive patients.17 In low-responsive individuals, ASA’s generally used dose of 100 mg/day may be insufficient for inhibiting platelet aggregation.17 Dose adjustments up to 200 mg/day may reduce the number of non-responders. The need for this adjustment may be related to different risk factors for atherosclerotic disease.17 In addition, treatment with 100 mg/day of ASA and 1000 mg/day of omega-3 fatty acid can reduce the number of non-responders to classical treatment and avoid undesirable effects such as bleeding and gastrointestinal alterations.17

Thus, the objective of this study was to evaluate the use of 200 mg/day of ASA or 100 mg/day of ASA combined with 1 g of omega-3 fatty acid in the response of patients with chronic coronary artery disease refractory to previous use of 100 mg/day of ASA and to determine polymorphisms of SNPs associated with treatment resistance. For this purpose, pharmacological analyses were performed, including in vitro platelet aggregation studies and genetic polymorphism analyses.

**Methods**

This is an analytical, experimental, and prospective study conducted between May 2013 and October 2015 in 152 patients screened at the outpatient clinic of the Dante Fazzanese Cardiology Institute (Fig. 1). This sample included individuals of both sexes, ages 31–81, with chronic coronary artery disease and under treatment with ASA at a dose of 100 mg/day for more than 30 days (Fig. 2). All participants signed an Informed Consent Form previously approved by the Research Ethics Committee of the Dante Fazzanese Cardiology Institute. The same committee approved this study under registration number CAAE:22534113.0.0000.5462. The authors are responsible for the fidelity of the study, the accuracy of the analyses, and data completeness.

**Experimental design**

In a first clinical evaluation (Time A) of participants under 10–12 h of fasting and 24 h after ingestion of the 100 mg/day dose of ASA (group ASA100), samples were collected for laboratory tests. Subsequently, patients were randomly divided into two treatment groups showing clinical and demographic data distributed homogeneously and without statistically significant differences (Table 1 of the supplementary data). Both groups underwent 30 days of treatment with the adjusted dosage. The ASA200 group received a dosage of 200 mg/day of ASA, and the ASA10003 group received 100 mg of ASA and 1 g of omega-3 fatty acid per day. At the end of the 30-day treatment and 24 h after administration of the last dose of the corresponding drugs, the second evaluation (Time B) of the patients was performed. In this clinical evaluation, samples were also taken for a new round of laboratory analyses.

**Biological samples and laboratory analyses**

Blood collection for biochemical and DNA analyses included the use of four 10 ml dry tubes, four 10 ml tubes with ethylene-diaminetetraacetic acid, two 10 ml tubes containing citrate, and one 10 ml tube containing fluoride. In addition, one 0.5 ml tube containing hinudin was used for blood collection from each patient for platelet aggregation analysis on the Multiplate platform (Roche, Switzerland). A 5 ml vial containing the first-morning urine, or from minimum retention of four hours, was also collected for urinalysis. Serum and nucleic acid voucher samples were stored in the Molecular Biology Laboratory for further studies after approval by the Ethics and Research Committee and with the patient’s authorization.
Biochemical tests using enzymatic-colorimetric and immunochemical methods were performed at the Clinical Laboratory of the Dante Fazzanese Institute. Thus, glyceria, total cholesterol as well as high-density lipoprotein (HDL), low-density lipoprotein (LDL) and very low-density lipoprotein (VLDL) fractions, triacylglycerides, complete blood count, serum urea, serum creatinine, serum uric acid, aspartate transaminase, alanine transaminase, fibrin degradation products (Dimer), high-sensitive C-reactive protein and Urine I (Ta 1) were determined.

The platelet function of the samples from the ASA100, ASA200, and ASA10003 groups was determined using the Multiplate analyzer platform (Roche, Switzerland). Analysis of cyclooxygenase-1 (COX-1) inhibition by ASA was performed using ASPItest (Roche, Switzerland), a reagent containing the COX-1-specific agonist arachidonic acid. Platelet function was also assessed using TRAPtest (Roche, Switzerland), which contains the thrombin's receptor activator peptide-6 (TRAP-6), a potent platelet activator. These analyses were performed according to the manufacturer's instructions. The results of the ASPItest at Time A were used to classify patients into ASA responders and non-responders (Table 2 of the supplementary data). Thus, patients were considered “highly responsive” up to 30 aggregation units (AU), “responders” from 30 to 40 AU, and “non-responders” above 40 AU.

**Plasma determination of thromboxane B2**

Dosage level of thromboxane B2 (TXB2) was performed on plasma samples from whole peripheral blood collected in an ethylenediaminetetraacetic acid-containing tube. TXB2 concentrations were determined using the Cusabio Human Thromboxane B2 ELISA kit according to the manufacturer's instructions. The quantification of TXB2 was performed by spectrophotometry, by reading absorbance at a wavelength of 450 nm, using the Multiskan GO platform (Thermo Fisher Scientific, United States). The readings were performed in duplicate and monitored using the standard curve provided by the manufacturer.

**Analysis of DNA polymorphisms**

The SNPs polymorphisms for the C/T substitution in the rs3842787 and rs3842798 regions of the PTGS1 gene of the COX-1 enzyme and the SNP for the T/C substitution in the rs5918 region of the ITGB3 gene of the glycoprotein IIbα (GPIIbα) subunit of the fibrinogen receptor were analyzed. These SNPs were chosen based on the information available in the NCBI's dbSNP database.18

Genomic DNA for these analyses was extracted from whole peripheral blood using the QIAamp DNA Blood Max Kit (QIA-GEN, GmbH, Germany) and the QIAcube automated system (QIA-GEN, GmbH, Germany). From these samples, the polymorphic regions were amplified by polymerase chain reaction (PCR) using the primer pairs 5'-CAGAGGAATGAAGGCCAG-3'/5'-CCCCATCCACCAAAAGACT-3' (PTGS1 rs3842787 region of 497 bp), 5'-GGGGATAGAAGCCAAAATAGGG-3'/5'-CTATGAGGATGAGCAGGTC-3' (PTGS1 rs3842798 region of 496 bp), and 5'-TGCTCCACATGGTAGGTAA-3'/5'-TCCCAAGACTTCCCCTGTA- (ITGB3 rs5918 region of 396 bp). The three primer pairs were selected using the Primer3 program v.0.4.019 and sequences available in the NCBI18 and Ensembl20 databases. The amplification reactions of the genomic DNA extraction samples were prepared using 1U
of Taq DNA polymerase (DNA Express Biotechnology, Brazil), 1× PCR buffer (DNA Express Biotechnology, Brazil), 200 μM of each dNTP (GE Healthcare, England), and 100 nM of each primer (DNA Express Biotechnology, Brazil), and conducted in a Veriti thermocycler (Applied Biosystems, United States). The PCR conditions consisted of an initial denaturation at 95 °C for 3 min, followed by 40 cycles of denaturation at 95 °C for 30 s, hybridization at 62 °C for 30 s, and extension at 72 °C for 1 min, with an extension at 72 °C for 10 min.

PCR products were analyzed by horizontal electrophoresis in a 1% agarose gel immersed in 0.5× TBE buffer (90 mM Tris–HCl, 90 mM boric acid, 2.0 mM EDTA, pH 8.0) for 30 min at 100 V and 60 mA. DNA was visualized under ultraviolet light after incorporating the intercalating agent GelRed (Biotium Inc., Hayward, United States). PCR products that showed fragments of expected size with an absence of nonspecific amplification were purified using the Illustra GFX PCR DNA and Gel Band kit (GE Healthcare, England). After treatment with the BigDye Terminator v3.1 Cycle Sequencing kit (Applied Biosystems), these purifications were sent to the Human Genome Study Center (Instituto de Biociências, USP, Brazil) for sequencing on an ABI 3730 DNA Analyzer (Life Technologies – Applied Biosystems). Sequencing results were analyzed in the Sequencing Analysis 5.3.1 software, using Base Caller KB and BioEdit Sequence Alignment Editor software. Some of these samples from the 3 genotypes confirmed by Sanger sequencing were used as controls in real-time PCR (qPCR) reactions.

The qPCR analyses were performed on a RotorGene thermocycler (Qiagen GmbH, Hilden, Germany) using the TaqMan SNP Genotyping Assay kit (Life Technologies, United States) and two probes for each polymorphism. These two probes, homologous to each of the studied alleles, are marked with 6-carboxyfluorescein (FAM probe) and 6-carboxy-4,7, 2', 7'-tetrachlorofluorescein (VIC probe). VIC/FAM labeling was used to identify C/T substitution polymorphism in PTGS1 rs3842787 and PTGS1 rs3842798 genes, and FAM/VIC labeling for T/C substitution polymorphism in ITGB3 rs5918 gene. The amplification reactions of the genomic DNA samples were prepared using reagents from the TaqMan Genotyping MasterMix Kit (Life Technologies, United States) according to the manufacturer’s indications, 0.5 μl of Taqman SNP Genotyping Assay 40× (Life Technologies, United States), and DNase-free water to a final volume of 20 μl. The qPCR conditions consisted of a 10 min cycle at 95 °C for activation of the HotStar Taq DNA polymerase, followed by 50 cycles of 15 s at 95 °C for DNA denaturation and ending with 90 s at 60 °C for probe hybridization and DNA extension. The data were analyzed using the Rotor-Gene Q – Pure detection program (QIAGEN, GmbH, Germany). At the end of the reactions, approximately 10% of the samples were sequenced by the Sanger method for genotyping confirmation.

Statistical analysis

Before comparing the quantitative variables of the various analyses, the Kolmogorov–Smirnov test was used to assess whether the parameters followed a normal distribution. The parametric Student’s t-test was used for data with normal distribution. For the remaining data, the non-parametric Mann–Whitney or Wilcoxon tests were used. To compare the frequencies of the qualitative variables, the Chi-square test was performed. All statistical analyses were conducted using the SPSS 20 software (SPSS Inc., United States), and the significance level adopted was P < .05. Still, the Hardy–Weinberg equilibrium was calculated for the study population after genotyping the polymorphisms in the PTGS1 and ITGB3 genes.

Results

Analysis of platelet aggregation and thromboxane dosage

From the initial analysis of platelet function in the ASA100 group, of all 152 patients, 38 non-responders were detected (25.2%). Subsequent tests of platelet function and plasma TBX2 concentration, including the ASA200 and ASA10003 treatment groups, showed that the reduction of ASPtest values was more significant (P = .001) in the non-responders’ group (NR) than in the responders’ (R) group (Fig. 3). However, only after treatment do the ASPtest values fall to the levels of responders (<40 AU) in the ASA200 group (Fig. 4). In addition, when the R and NR groups were analyzed separately, it was possible to observe a correlation between platelet function test results and TBX2 in the R group of the ASA10003 treatment (P = .039) (Table 1). By organizing responder and non-responder data by treatment groups (Fig. 5), it can be noted that, although the treatment effect was not statistically different between groups, non-responders who used ASA200 for 30 days had ASPtest values reduced to the level of reclassification as responders (<40 AU), which did not occur in the ASA10003 group.

There was no statistically significant difference between treatment groups concerning platelet function (ASPtest and TRAPtest) and TBX2 production (Fig. 6). TRAPtest values were within the expected range (84–128), demonstrating that platelet function was preserved in these patients. There was also no correlation between ASPtest results and plasma TBX2 concentration (Table 1).

Polymorphisms

The rare allele of the PTGS1 rs3842787 polymorphism is associated with a worse ASPtest response. In comparison, patients
Fig. 4 – Effect of treatments, expressed as median and interquartile range values, on platelet function measured by ASPItest (ASP) and TRAPtest (TRAP), and plasma concentration of thromboxane B2 (TBX2) in the groups of responders (R) and non-responders (NR) to acetylsalicylic acid (ASA) at Times A (after 30 days of treatment with ASA 100 mg, ASA100) and B (after 30 days of treatment with ASA 200 mg, ASA200, or ASA 100 mg and omega-3, ASA100O3). a Mann–Whitney test. b Student’s t-test.

<table>
<thead>
<tr>
<th>Treatments</th>
<th>Non-responder</th>
<th>Responder</th>
</tr>
</thead>
<tbody>
<tr>
<td>ASP</td>
<td>TBX2</td>
<td>ASP</td>
</tr>
<tr>
<td>Spearman’s coefficient of correlation</td>
<td>1.000</td>
<td>0.051</td>
</tr>
<tr>
<td>P-value</td>
<td>.001</td>
<td>.030</td>
</tr>
<tr>
<td>N</td>
<td>150</td>
<td>57</td>
</tr>
</tbody>
</table>

a ASP: ASPItest.

b TBX2: thromboxane B2.

carrying the rare allele of the rs5918 polymorphism in the ITGB3 gene showed a worse response concerning TBX2 concentration after the 30-day treatment (Table 2). No statistical difference was found in the genotypic and allele frequencies of the studied polymorphisms between the R and NR groups and between the ASA200 and ASA100O3 treatments (Table 3 of the supplementary data). Therefore, the rare alleles were equally distributed between R and NR groups and treatment groups. The genotypic distribution of the studied population agreed with the predictions of the Hardy–Weinberg law (Table 4 of the supplementary data).

Biochemical analysis

There was a significant reduction in VLDL and triglyceride values in the ASA200 (P = .005 and P = .002, respectively) and ASA100O3 (P = .006 and P = .004, respectively) groups after 30 days of treatment. However, this reduction was not significantly different between both treatments. It was also possible to observe a significant reduction in the number of monocytes (P = .024) in the ASA100O3 treatment group compared to the ASA200 group.

Discussion

Patients initially classified as non-responders showed a significant reduction in ASPItest values in the analysis of platelet function and plasma TBX2 concentration after 30 days of ASA200 treatment. In this treatment group, ASPItest values dropped to < 40 AU, reaching the responder classification. This reduction was not detected in the ASA100O3 treatment group. It is important to note that the doses of ASA used in this
study were low, agreeing with other studies reporting the use of doses of 50, 100, 325, and 1000 mg.\textsuperscript{21,22}

Platelet function was preserved in patients in the ASA200 and ASA100O3 treatment groups, as demonstrated by TRAPtest values, which were within the expected range (84–128). Thus, there was no statistically significant difference in platelet function measured by the ASPtest and TRAPtest between the ASA200 and ASA100O3 treatment groups regarding the TBX2 production. Interestingly, patients in both groups had reduced VLDL and triglyceride values after 30 days of treatment, with no significant difference between ASA200 and ASA100O3 treatments. Also, in the ASA100O3 group, there was a decrease in monocytes, which may be related to the anti-inflammatory action of omega-3 on atherosclerotic plaque.

These results are encouraging in light of evidence from studies reporting a 30% reduction in inflammatory activity after four weeks of treatment with omega-3 fatty acid intake.\textsuperscript{23,24}

The correlation between the results of the platelet function test and TBX2 production of the responder’s group agrees with Kidson-Gerber et al.,\textsuperscript{25} which show that measuring TBX2 levels is a direct and easy-to-apply method to analyze the effect of ASA on platelets. Hence, this method has an adequate specificity in measuring the ability of platelets to produce thromboxane A2 and, consequently, directly measures the effect of ASA on platelet function.\textsuperscript{25} These platelet response analysis techniques are accessible to medical practice. In addition, our results show that monitoring the patient’s response to anticoagulant treatment is fundamental in clinical practice.
In this way, the physician can timely identify a non-responder patient, considering the techniques used in our study. Analysis of lack of response to ASA should not be limited to its pharmacological effect on COX-1 thromboxane-dependent inhibition. Treatment failure should also be analyzed concerning other events occurring under the use of ASA or when there is inhibition of platelet activation. Environmental factors, gender, obesity, smoking, drug interactions (NSAIDs), and conditions associated with increased platelet activity, such as exercise, dyslipidemia, systemic hypertension, and acute diseases, can modify the inhibitory action of ASA on platelets. Therefore, specific tests for the effect of ASA, such as those of TBX2 levels, are adequate for differentiating between an insufficient pharmacological action of ASA or other causes of treatment failures in non-responders. In patients who respond to ASA in TBX2 tests and on aggregometer platforms, ASA's inability to inhibit A2-dependent thromboxane production and high innate platelet reactivity may occur together, thus confirming, in these cases, a reduced effect of ASA in preventing cardiovascular events. Therefore, in clinical practice, it is helpful to use a test of the pharmacological effect of ASA (serum TBX2 level) and a platelet function test (Multiplate).

Regarding the polymorphisms of the PTGS 1 – (COX-1) gene, the rare allele rs3842787 was found to be associated with a worse response in ASPitest in non-responders. On the other hand, patients with the rare allele rs5918 of the ITGB3 3 gene (GPIIIa) have a worse response regarding the concentration of TBX2 after 30 days of treatment. Various candidate genes have been suggested to explain the genetic polymorphisms impacting the analysis of interindividual response variation to ASA. A meta-analysis of 31 studies, including 50 polymorphisms of 11 genes and 2834 patients, reported the polymorphism in the rs5918 region of the ITGB3 gene of the GPIIIa platelet receptor as the most frequently investigated and reported in association with ASA resistance. These authors showed no association of polymorphism in the rs3842787 region of the COX-1 gene (A842G/CSOT) with ASA resistance. Maree et al. genotyped 144 patients with coronary artery disease and, of the five polymorphisms studied, showed a significant effect of antithrombotic modulation of ASA for the C50T variant (rs3842787). Similar observations were made by Lepäntalo et al. Su et al. showed that the 807T (GPIIIa) – 155918 allele confers 3.8 times more resistance to the effect of ASA in 200 patients with atherosclerosis than in 807 homozygotes from a Chinese population.

ASA resistance analyzed by genetic variants considers that polymorphisms vary according to ethnicity. Thus, Wagner et al. detected higher resistance to ASA for Europeans with rs3842787 and Africans with rs3842792. In the Chinese population, gene polymorphism rs5918 strongly correlates with ASA resistance. More specifically, Shiffman et al. showed that carriers of an SNP related to plasma lipoprotein levels benefit more from treatment with ASA than non-carriers. Studies like these may lead to a personalized genetic profile that more accurately identifies which patients will benefit most from using ASA. Despite their usefulness, the cost of genetic analyses may limit their application in particular clinical practices. However, these analyses are increasingly widespread and accessible to public and private health centers. Therefore, genetic analyses can become valuable tools to analyze SNP polymorphisms in non-responders and thus guide the most appropriate treatment for each case.

The efficacy and safety of ASA have been extensively analyzed in apparently normal populations at low, moderate, or high risk for cardiocirculatory events and populations with

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Table 2 – Effect of rare alleles on ASPitest results and concentration of plasma thromboxane B2.

<table>
<thead>
<tr>
<th>Polymorphism</th>
<th>Test</th>
<th>CC genotype</th>
<th>CT+TT genotype</th>
<th>P*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td>Median (25%; 75%)</td>
<td>Median (25%; 75%)</td>
<td></td>
</tr>
<tr>
<td>PTGS1 rs3842798 C&gt;T</td>
<td>ASPitest_A (AU)</td>
<td>18.5 (8.5; 59)</td>
<td>23 (15.8; 42.5)</td>
<td>.492</td>
</tr>
<tr>
<td></td>
<td>ASPitest_B (AU)</td>
<td>14.5 (9.3; 27)</td>
<td>22 (15; 36.5)</td>
<td>.085</td>
</tr>
<tr>
<td></td>
<td>ASPitest_delta (AU)</td>
<td>−9.0 (−38.8; 2.2)</td>
<td>−2.0 (−5.5; 7.5)</td>
<td>.214</td>
</tr>
<tr>
<td></td>
<td>TBX2_A (pg/ml)</td>
<td>128.8 (38.1; −)</td>
<td>163.6 (84.5; −)</td>
<td>.775</td>
</tr>
<tr>
<td></td>
<td>TBX2_B (pg/ml)</td>
<td>167.9 (91.5; −)</td>
<td>116.6 (60.9; −)</td>
<td>.335</td>
</tr>
<tr>
<td></td>
<td>TBX2_delta (pg/ml)</td>
<td>129.8 (−276.3; −)</td>
<td>−53.4 (−112; −)</td>
<td>.282</td>
</tr>
<tr>
<td>PTGS1 rs3842787 C&gt;T</td>
<td>ASPitest_A</td>
<td>25 (17; 51)</td>
<td>18 (12.5; 22)</td>
<td>.023*</td>
</tr>
<tr>
<td></td>
<td>ASPitest_B</td>
<td>22 (13; 36)</td>
<td>21 (17; 33)</td>
<td>.810</td>
</tr>
<tr>
<td></td>
<td>ASPitest_delta</td>
<td>−3.0 (−18; 5)</td>
<td>4.0 (0.25; 11.75)</td>
<td>.028</td>
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<td></td>
<td>TBX2_A</td>
<td>170.7 (82.3; 292.5)</td>
<td>100.1 (62.9; 140.9)</td>
<td>.108</td>
</tr>
<tr>
<td></td>
<td>TBX2_B</td>
<td>116.6 (65.2; 199.9)</td>
<td>127.9 (53.6; 202.3)</td>
<td>.693</td>
</tr>
<tr>
<td></td>
<td>TBX2_delta</td>
<td>−44.5 (−120.9; 37.9)</td>
<td>27.8 (−13.7; 65.8)</td>
<td>.114</td>
</tr>
<tr>
<td>ITGB3 rs5918 T&gt;C</td>
<td>ASPitest_A</td>
<td>20 (15.75; 34.5)</td>
<td>23.5 (15; 55.25)</td>
<td>.446</td>
</tr>
<tr>
<td></td>
<td>ASPitest_B</td>
<td>21.5 (12.75; 42.75)</td>
<td>22 (14.5; 32)</td>
<td>.792</td>
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<td></td>
<td>ASPitest_delta</td>
<td>0 (−15; 8)</td>
<td>−2.0 (−18; 6)</td>
<td>.388</td>
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<td></td>
<td>TBX2_A</td>
<td>224.3 (92; 327.4)</td>
<td>162.0 (81.3; 162)</td>
<td>.181</td>
</tr>
<tr>
<td></td>
<td>TBX2_B</td>
<td>90.6 (69.9; 90.6)</td>
<td>128.3 (58.3; 128.3)</td>
<td>.418</td>
</tr>
<tr>
<td></td>
<td>TBX2_delta</td>
<td>−118.9 (−247.8; −11.8)</td>
<td>−4.6 (−72.8; 86)</td>
<td>.013*</td>
</tr>
</tbody>
</table>

A: response to treatment ASA100 (first visit); B: response to treatment ASA200 or ASA100003 (second visit); delta: B-A. C: cytosine; T: thymine; CC: cytosine homozygous genotype; TT: thymine homozygous genotype; CT: heterozygous genotype.

* P-value from Mann–Whitney test comparisons.

a Statistically significant, p < .05; TBX2: thromboxane B2.
diseases with or without a history of AMI and/or stroke.\textsuperscript{37-39} Specifically, a meta-analysis on primary atherosclerosis prevention studies showed that ASA reduced the total number of cardiovascular events by 10%, primarily by reducing non-fatal AMI (20%).\textsuperscript{40} However, the reduction in mortality from cardiovascular causes and neoplasms was negligible, 1% and 7%, respectively. Bleeding occurred in 30% of individuals using ASA. These “modest benefits” and the high risk of bleeding do not justify the routine use of ASA in the primary prevention of cardiovascular diseases. Hence, there is a need to identify population groups that will benefit from treatment and establish adequate doses, aiming at therapeutic effectiveness and low risk of bleeding. In our study, patients with the s38478 allele had no higher incidence of thrombosis or bleeding. Still, none of these clinical signs were detected or reported by patients from both treatment groups during clinical evaluation at Time B. However, new studies will be developed to analyze these clinical signs in more detail.

**Limitations**

Some limitations of our study are related to a relatively small sample size, the usage of 1 g of omega-3, despite literature describing 1–4 g per day, and the lack of in vitro analysis of platelet aggregation on diabetic patients that were not the subject of this study. In addition, adherence to medication intake was monitored during clinical evaluations and by telephone. Therefore, our results should be carefully interpreted since they originated from a single-center study on a limited number of patients. This limited n may influence the results of statistical analyses. However, our results show helpful information for clinical practice and the design of more complex, multi-center studies on more patients. Among the studies derived from our research, we can include the analysis of temporal variation in platelet activity, stratified studies in responders and non-responders, or analyses of isolated non-responders. These future studies may apply our methodology to assess, for example, the response of patients with atrial fibrillation to oral anticoagulant administration.

**Conclusions**

Combining platelet function analysis and genetic testing is key to identifying and classifying ASA non-responder patients. In this way, the best available treatment can be administered to each patient. Within these possibilities, we recommend considering ASA200 treatment for 30 days to reduce the ASPItest values of non-responder patients down to the responder level. Interestingly, the use of the two treatments in the present study showed no statistically significant difference between the two treated groups regarding absence of hematuria and bleeding, constituting safe treatment alternatives. In addition, treatment with 1 g/day of omega-3 fatty acid, one of the lowest doses concerning those reported in the literature,\textsuperscript{41} reduced the number of monocytes. The detection of the rs384787 alleles associated with the worst response to ASPItest and the rs5918 allele to the worst response in the TBX2 test, combined with the therapeutic findings described, reinforces the usefulness of pharmacogenetics, enabling personalized patient treatment, and avoiding the undesirable effects of gastrointestinal symptoms and bleeding of the ASA.

**What is known about the topic?**

Antiplatelet agents like ASA have a central function in the prevention of atherothrombosis. However, some patients using ASA continue to suffer atherothrombotic events, and approximately 20% will have some recurrent vascular event. Some of these side effects, such as bleeding, raising the debate about creating a specific therapy for each patient. In addition, the combined use of other drugs, such as omega-3, can potentiate the antiaggregant effects of ASA without its side effects.

**Does it contribute anything new?**

We show that patients initially classified as non-responders obtained a significant reduction in ASPItest values and plasma TBX2 concentration after 30 days of ASA200 treatment. In addition, there is a significant reduction in VLDL and triglyceride values in the ASA200 and ASA10003 treatments. The absence of hematuria and bleeding confirms both treatments as safe therapeutic alternatives. We detected rare alleles associated with worst responses to ASPItest and TBX2 concentration analysis.

**Acknowledgments**

We thank the Adib Jatene Foundation for the financial support to conduct genetic and platelet aggregation analyses and the Associação Fundo de Incentivo a Pesquisa for collaborating in the biochemical tests. We also thank two anonymous reviewers for their valuable comments on our manuscript.

**Funding**

This work received support from the Adib Jatene Foundation, Av. Dr. Dante Pazzanese, 500 – Vila Mariana, São Paulo, Brasil – CEP 04014-002.

**Authors’ contributions**

Each author contributed significantly to the submitted work regarding the research conception (M.M. Cartocci, A. Guerra de Moraes Rego Sousa, and D. Armaganijan), data analyses (M.M. Cartocci and M.H. Hirata), and writing of the manuscript (M.M. Cartocci, A. Guerra de Moraes Rego Sousa, and D. Armaganijan, M. Batlouni, and M.H. Hirata).

**Conflicts of interest**

None.
Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at doi:10.1016/j.rcl.2023.07.002

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