

Efficacy of piroxicam for postoperative pain after lower third molar surgery associated with *CYP2C8*3* and *CYP2C9*

Adriana Maria Calvo
Paulo Zupelari-Gonçalves
Thiago José Dionísio
Daniel Thomas Brozoski
Flávio Augusto Faria
Carlos Ferreira Santos

Department of Biological Sciences,
Bauru School of Dentistry, University
of São Paulo, São Paulo, Brazil

Objective: Nonsteroidal anti-inflammatory drugs (NSAIDs) are metabolized by the cytochrome P450 enzymes (CYPs), predominantly *CYP2C8* and *CYP2C9*. The aim of this study was to evaluate the possible association of polymorphisms in the *CYP2C8*3* and *CYP2C9* genes with the clinical efficacy of oral piroxicam (20 mg daily for 4 days) after lower third molar surgeries with regard to postoperative pain, swelling, trismus, adverse reactions, need for rescue medication and the volunteer's overall satisfaction.

Materials and methods: For this purpose, 102 volunteers were genotyped for *CYP2C8*3* and *CYP2C9* polymorphisms. Briefly, genomic DNA was isolated from saliva collected from volunteers subjected to invasive lower third molar surgeries, and the preoperative, intraoperative and postoperative parameters were collected and analyzed.

Results: An equal amount of piroxicam sufficiently managed postoperative pain and inflammatory symptoms, with visual analog pain scores typically <40 mm for all genotypes investigated. Furthermore, only two out of 102 volunteers heterozygous for *CYP2C8*3* and *CYP2C9*3* reported adverse side effects.

Conclusion: In general, slow metabolizers of piroxicam, who were volunteers with mutant alleles, were indifferent from normal metabolizers with the wild-type alleles and therefore did not require specialized piroxicam doses to manage postoperative pain and inflammation.

Keywords: piroxicam, lower third molar surgery, P450, *CYP2C8*, *CYP2C9*, pharmacogenetics

Introduction

The variability in each individual's response to the same dose of a drug is a major cause of side effects. In many cases, this variability is connected to polymorphisms in genes encoding enzymes responsible for metabolizing drugs.¹ In particular, carriers of mutations in genes encoding drug-metabolizing enzymes tend to have higher plasma levels of the drug^{2,3} and an increase in the frequencies and severity of adverse reactions, when compared to nonmutants, when taking the usual doses of a drug that is metabolized by the affected enzyme.⁴⁻⁶ In other words, these mutations, which may consist of an alteration in one nucleotide, can cause absent, reduced or increased enzymatic activity.^{2,7} In particular, cytochrome P450 2C8 (*CYP2C8*) and 2C9 (*CYP2C9*) belong to one of the major families of enzymes involved in drug metabolism.

The genes encoding *CYP2C8* and *CYP2C9* are grouped in two consecutive clusters on chromosome 10.³ Briefly, a single-nucleotide polymorphism (SNP) is a variant DNA sequence wherein one nucleotide differs between common populations in a species.⁸ Furthermore, these SNPs are commonly found in exons and introns. The common nomenclature for the normal wild-type (wt) SNP for *CYP2C9* is noted with

Correspondence: Carlos Ferreira Santos
Department of Biological Sciences, Bauru
School of Dentistry, University of São
Paulo, Alameda Doutor Octávio Pinheiro
Brisolla, 9-75, Bauru, São Paulo
17012-901, Brazil
Tel +55 14 3235 8295
Email cfsantos@fob.usp.br

a *1; two common polymorphic variants are allele *2, with the amino acid change R144C, and allele *3 with the amino acid change I359L; each individual commonly carries two alleles and may be either homozygous or heterozygous, in both cases, mutant (mt).⁹

Both CYP2C8 and CYP2C9 enzymes are associated with the metabolism of several drugs used clinically, some of which possess narrow therapeutic margins.¹⁰ The clinical importance of CYP2C8 and CYP2C9 polymorphisms with regard to nonsteroidal anti-inflammatory drugs (NSAIDs) mainly involves two factors: 1) these polymorphisms are often associated with an increased number of adverse reactions; and 2) these adverse reactions are often more serious. Notably, “slow” metabolizers of piroxicam (eg, people with CYP2C9*1/*3 and *3/*3 alleles), when compared to normal metabolizers (eg, people with CYP2C9*1/*1 wt alleles), might respond differently to the same dose of piroxicam, although it remains unknown whether piroxicam is metabolized differently by either CYP variant depending on whether these molecules are in ketonic or enolic tautomeric forms.

Data regarding the main role of CYP2C8 in the metabolism of piroxicam are inconsistent, but some studies suggest that SNPs of CYP2C8 tend to decrease the clearance of NSAIDs in general.⁹ Several studies have associated the genotypes of CYP2C8 and CYP2C9 with the pharmacokinetics and pharmacodynamics of piroxicam and other NSAIDs in healthy individuals.^{4,11–15} In contrast, this study aimed to evaluate the possible association of polymorphisms of CYP2C8*3, CYP2C9*2 and CYP2C9*3 with the clinical efficacy of oral piroxicam (20 mg daily for 4 days) after invasive lower third molar surgeries with regard to pain, swelling, trismus, need for rescue medication, volunteer satisfaction and adverse reactions to piroxicam. Briefly, the volunteers selected for this study required invasive surgeries for lower third molars with the following classifications: Pell and Gregory class IIB, IIC, IIIA, IIIB and IIIC and/or Winter horizontal, distal, inverted and transalveolar.¹⁶ It was hypothesized that postoperative pain would be effectively managed by a daily 20 mg oral dose of piroxicam for all genotypes, but that volunteers with mutant CYP2C8*3 and CYP2C9 genotypes would report less postoperative pain after lower third molar surgery.

Materials and methods

Registration and study design

The current study was performed in accordance with the Declaration of Helsinki and was approved by the ethics committee of Bauru School of Dentistry, University of São

Paulo, Brazil National Research Ethics System (CAAE number: 20657913.7.0000.5417) in accordance with resolution 196/96 of the National Council of Health/Ministry of Health. This study was registered with Universal Trial (U1111-1155-3545); ClinicalTrials.gov (NCT02450487). All volunteers provided signed informed consent agreements after receiving all instructions and before undergoing any interventions.

Briefly, 105 volunteers (≥18 years old) visiting the Clinical Physiology and Pharmacology Laboratory (LAFFIC) requiring the extraction of at least one lower third molar that was included and/or impacted (as determined by panoramic radiography) participated in this study. Three volunteers did not return for postoperative measurements and were completely excluded from this study (Figure 1). Saliva samples were collected to genotype all volunteers, and postoperative data were used to judge the clinical efficacy of piroxicam (20 mg daily for 4 days) after lower third molar surgeries.

Eligibility criteria included adults without inflammation, infection and systemic diseases that could possibly interfere with the study. Exclusion criteria were as follows: any history of allergy, gastrointestinal bleeding/ulcers, kidney disease, asthma or any known allergic sensitivity to any NSAID. Pregnant women were also excluded from the study, as well as volunteers who used antidepressants, diuretics or NSAIDs 7 days prior to the start of the experiments. All surgeries were performed between December 2013 and October 2014, after ethical approval. The sample size calculation for the current study was based on the facial edema and trismus parameters from our previous research article.¹⁷ Taking into account the facial edema data with 94% power and effect size of 0.7, the number of patients required was 102; considering trismus with 80% power and effect size of 0.56, the number of patients remained 102.

All surgeries required bone removal, and, in some cases, tooth section. The positions of molars were determined with panoramic radiographs. No antibiotics were prescribed during the study to avoid possible bias in the drug protocol.¹⁷

Surgery intervention and assessments

All surgeries were performed by the same maxillofacial surgeon (PZG). One cartridge (1.8 mL) of local anesthetic (4% articaine with 1:200,000 epinephrine) was injected near the buccal, lingual and inferior alveolar nerves to achieve truncal anesthesia. After 3 minutes and when the patient confirmed a loss of sensation of the lower lip, a half cartridge (0.9 mL) of local anesthetic was injected to obtain complete and consistent anesthesia. A half cartridge (0.9 mL)

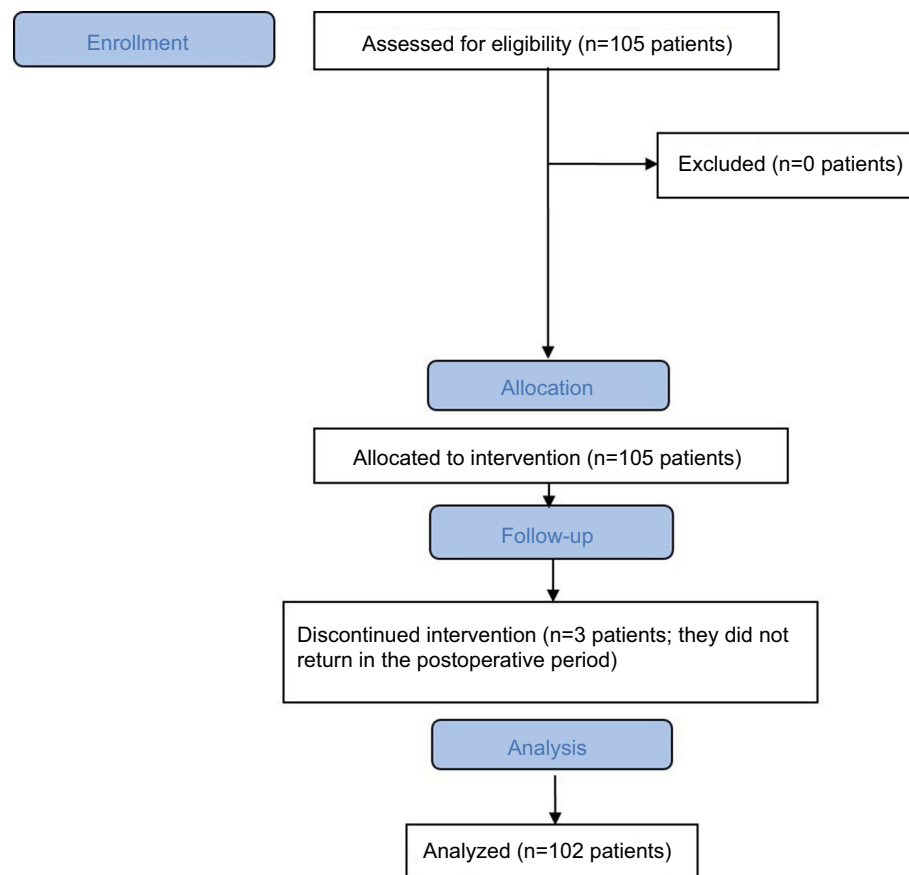


Figure 1 CONSORT flowchart.

of local anesthetic was also administered using the terminal infiltration technique to reduce bleeding and ensure mucosal anesthesia.^{16–24}

The protocol used for postoperative pain medication after extraction of the third lower molars was a self-administered 20 mg tablet of piroxicam once every 24 hours for 4 days. Oral rescue analgesic medication was also available as needed throughout the study; for that, 750 mg tablets of paracetamol were provided to all volunteers, who were instructed to record the time of consumption while maintaining the use of piroxicam.^{16–19,21–26}

The preoperative, intraoperative and postoperative parameters evaluated are reported in Table 1. Briefly, postoperative mouth opening (millimeters) between the mesial–incisal corners of the upper and lower right central incisors during maximum opening of the jaws was measured and recorded before surgery as well as during the second and seventh postoperative days, according to Ustun et al.^{16,24,27}

For swelling measurements of the facial edema, methods were used according to Ustun et al, which sums the following measures: 1) the distance between the lateral canthal ligament and the gonion; 2) the distance between the tragus

and the labial commissure of the mouth; and 3) the distance between the tragus and the soft tissue of pogonion.^{16,24,27} The preoperative sum of these three measures was considered the baseline. The percentage difference between the baseline value and the values of the second and seventh postoperative periods for swelling indicated the facial edema.

Genotyping *CYP2C8*3* and *CYP2C9*

All volunteers were instructed to expectorate up to 4 mL of unstimulated saliva into a provided sterile polypropylene collection tube. Approximately 3 mL of the tube's contents were then placed in a sterile microcentrifuge tube using a sterile pipette and immediately stored in a freezer at -20°C until the genomic DNA was extracted. Finally, a QIAamp DNA Mini Kit (catalog number 51104; Qiagen®, Hilden, Germany) was used to extract genomic DNA from the saliva.

Genomic DNA was genotyped using the Taqman® system for performing real-time polymerase chain reaction (qPCR) using a ViiA™ 7 system (Applied Biosystems®, Waltham, MA, USA). Each reaction was performed in duplicate. Specifically, manufactured assays were used and validated by Applied Biosystems® with the following catalog numbers:

Table 1 Study parameters evaluated

Parameters	Units
Total volume of local anesthetic	Milliliters
Onset of anesthetic agent action	Minutes
Surgery duration	Minutes
Overall experience of surgery reported by volunteer	5-point scale: 1, poor; 2, fair; 3, good; 4, very good; 5, excellent
Quality of anesthesia	3-point scale: 1, no discomfort during surgery; 2, any discomfort without anesthesia required; 3, any discomfort with anesthesia required
Surgery difficulty	3-point scale: 1, no need for osteotomies without tooth sectioning; 2, need for osteotomies without tooth sectioning; 3, need for osteotomies as well as tooth sectioning complicated
Intraoperative bleeding	3-point scale: 1, minimal bleeding; 2, normal bleeding; 3, excessive bleeding
Quality of wound healing	7th day, 3-point scale: 1, normal healing without inflammation; 2, delayed healing; 3, healing complicated by inflammation or local infection with or without purulent material
Adverse reactions	Observed by the surgeon or reported by the volunteer, during the surgery in the first postoperative hour and during the 2nd and 7th days after surgery
Systolic, diastolic and mean arterial pressure; heart rate; and oxygen saturation ^a	Millimeters of mercury (mmHg); beats per minute (bpm); and percentage saturation of peripheral oxygen (% SpO ₂)
Body temperature	Preoperative period, 2nd, 7th postoperative day (degrees centigrade, °C)
Mouth opening	Preoperative period, 2nd, 7th postoperative day (millimeters)
Facial swelling	Preoperative period, 2nd, 7th postoperative day (millimeters)
Subjective evaluation of postoperative pain	VAS (0–100 mm)
Total amount of rescue medication	Number of tablets

Note: ^aDixtal[®] (model DX2021; Dixtal Biomédica Industria and Comércio Ltda, Marília, São Paulo, Brazil; ANVISA/MS 10293490035, model number 101503732).

Abbreviation: VAS, visual analog scale.

C_25628505_10, C_27104892_10, C_25625794_10 and C_256257782_20; *CYP2C9* [*CYP2C9**2-rs1799853 (C430→T) and *CYP2C9**3-rs1057910 (A1075→C)] and *CYP2C8* [*CYP2C8**3-rs11572080 (G416→A) and rs10509681 (A1196→G), respectively]. DNA samples were quantified and qualified using TaqMan[®] Genotyping Master Mix (catalog number: 4381656; Applied Biosystems[®]) to determine the *CYP* genotypes.

Statistical analysis

The G*Power software v.3.0.10 was used for the sample calculation. Data were analyzed using Microsoft[®] Excel 2002 (version 10.6871.6870) and GraphPad Prism (version 4.0). Briefly, data were tested for normality using the Shapiro–Wilk test. When data were normally distributed, comparisons among and between groups were made using the unpaired *t*-test. For all nonnormally distributed data, statistical comparisons between two independent groups were performed using the Mann–Whitney *U* test, while the Kruskal–Wallis test by ranks was used for analyzing nonparametric data with more than two groups. Lastly, binary data (eg, the absence or presence of an adverse reaction) were compared using Fisher's exact test.

Statistical significance was set at 0.05. Normally distributed data are reported as mean ± SD, whereas nonnormally distributed data are reported as median with the interquartile range (IQR).

Results

Table 2 reports the demographic distribution and intraoperative parameters of the 102 volunteers with lower third molar surgeries. Human subjects were between the ages of 14 and 58 years and came from the state of São Paulo, Brazil.

Excluding the transalveolar position, the third molar position of the tooth to be extracted was evenly distributed ($P=0.61$). The transalveolar-positioned third molars overall were significantly less common (2.9%) than other molar positions (mesioangular, vertical, distoangular or horizontal) when compared to positions being equally distributed (chi-square test, $P=0.00036$).

Hemodynamic parameters obtained during surgery fluctuated slightly during the different phases of extraction, yet no unusual peaks in blood pressure, heart rate and oxygen saturation were observed (data not shown). Additionally, during and immediately following surgery, no adverse reactions were found in any of the volunteers. Throughout the entire study, only two volunteers reported adverse side effects; on the second day, one volunteer reported sleepiness (*CYP2C8**3 mt and *CYP2C9**1/*3 genotype) and one volunteer (*CYP2C8**3 mt and *CYP2C9**1/*3 genotype) reported stomachaches. In addition, as indicated by some of the postoperative parameters, a volunteer's allele for *CYP2C8**3 or *CYP2C9* was not correlated with any variation in the anti-inflammatory effects of piroxicam.

Table 2 Preoperative, intraoperative and postoperative parameters

Parameters	All (n=102)	
	Median	IQR
Age, years	24	7
Lower third molar position	n	%
Distoangular	27	26
Horizontal	27	26
Mesioangular	19	19
Transalveolar	1	1
Vertical	28	27
Surgery	Median	IQR
Duration, minutes	10	7
Difficulty (score assessed by surgeon)	3.0	1.0
Intraoperative bleeding (score assessed by surgeon)	1.0	0.1
Quality of surgery (score assessed by volunteer)	3.5	1.0
Quality of wound healing (score assessed by surgeon)	1.0	0.0
Local anesthetic, 4% articaine with 1:200,000 epinephrine	Median	IQR
Quantity, mL	2.7	0.9
Onset, minutes	1.0	0.5
Quality (score assessed by surgeon)	1.5	2.0
Mouth opening	Average	SD
Preoperative measurement, mm	47	7
	Median	IQR
2-day postoperative measurement, percentage of preoperative	42	18
7-day postoperative measurement, percentage of preoperative	71	31
Facial swelling	Median	IQR
Preoperative measurement, mm	37	4
2-day postoperative measurement, percentage of preoperative	104	4
7-day postoperative measurement, percentage of preoperative	100	2
Underarm temperature, °C	Median	IQR
Preoperative measurement	36.2	0.7
2-day postoperative measurement	36.3	0.7
7-day postoperative measurement	36.1	0.6
Paracetamol rescue medication	Median	IQR
Time between first consumption and surgery, hours	3.5	3.5
	Average	SD
VAS during first consumption, mm	53	25
	Median	IQR
Total quantity consumed, mg	2250	3750

Abbreviations: IQR, interquartile range; VAS, visual analog scale.

The distribution of genotypes is reported in Table 3. Overall, all the volunteers grouped together were 72.5% homozygous for the wt *CYP2C8*3* allele and 27.5% heterozygous or homozygous for the mt *CYP2C8*3* allele (Table 3). Furthermore, volunteers grouped together were most commonly homozygous for the wt *CYP2C9*1/*1* allele (66.7%), followed by the mt variants **1/*2* (19.6%), **1/*3* (7.8%) and **3/*3* (5.9%), with no **2/*2* or **2/*3* genotypes found in the study population (Table 3). Table 4 reports the frequencies of the *CYP2C8*3* alleles compared to the *CYP2C9* alleles found in the population of volunteers.

Piroxicam was effective in the control of pain regardless of the *CYP* haplotype (*CYP2C8*3* and *CYP2C9*) carried by volunteers (Figure 2). There was a trend of lower levels of reporting pain (visual analog scale [VAS]) in volunteers carrying the *CYP2C9*1/*1* wt allele, though a statistically significant difference was not found between groups. This trend is justified considering reports in the literature that ancient homozygous individuals for *CYP2C9* are normal metabolizers of piroxicam.

It is noted that the average pain scores reported by the volunteers is <44 mm, which demonstrates a very high pain control profile for piroxicam independent of the *CYP* haplotype carried by the volunteers.

Discussion

This is the first report focusing on the influence of pharmacogenetics of the P450 cytochrome family (*CYP2C8*3* and *CYP2C9*) on the control of postoperative inflammatory signs after minor oral surgery. The vast majority of articles are related to pharmacokinetics, metabolism and adverse reactions of NSAIDs, such as gastrointestinal bleeding,^{2-6,10,12,13,28}

Table 3 Genotype frequencies of *CYP2C8*3* and *CYP2C9*

Allele	All (n=120)	
	n	%
<i>CYP2C8*3</i>		
Wild type (wt)	74	72.5
Mutant (mt)	28	27.5
<i>CYP2C9</i>		
<i>*1/*1</i>	68	66.7
<i>*1/*2</i>	20	19.6
<i>*2/*2</i>	0	0.0
<i>*2/*3</i>	0	0.0
<i>*1/*3</i>	8	7.8
<i>*3/*3</i>	6	5.9

Abbreviation: CYP, cytochrome P450.

Table 4 Genotypic frequencies of *CYP2C8* relative to *CYP2C9* in a Brazilian population of volunteers with third molar surgeries

Genotypes <i>CYP2C9</i>	<i>CYP2C8</i> *3						All	
	*1/*1		*1/*3		*3/*3			
	<i>rs10509681</i>	<i>rs11572080</i>	<i>rs10509681</i>	<i>rs11572080</i>	<i>rs10509681</i>	<i>rs11572080</i>	<i>rs10509681</i>	<i>rs11572080</i>
*1/*1	63 (61.8%)	63 (61.8%)	5 (4.9%)	5 (4.9%)	0 (0.0%)	0 (0.0%)	68 (66.7%)	68 (66.7%)
*1/*2	2 (2.0%)	4 (3.9%)	17 (16.7%)	16 (15.7%)	1 (1.0%)	0 (0.0%)	20 (19.6%)	20 (19.6%)
*2/*2	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)
*2/*3	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)	0 (0.0%)
*1/*3	6 (5.9%)	7 (6.9%)	2 (2.0%)	1 (1.0%)	0 (0.0%)	0 (0.0%)	8 (7.8%)	8 (7.8%)
*3/*3	4 (3.9%)	4 (3.9%)	2 (2.0%)	2 (2.0%)	0 (0.0%)	0 (0.0%)	6 (5.9%)	6 (5.9%)
All	75 (73.5%)		26 (25.5%)		1 (1.0%)		102 (100%)	

Abbreviation: CYP, cytochrome P450.

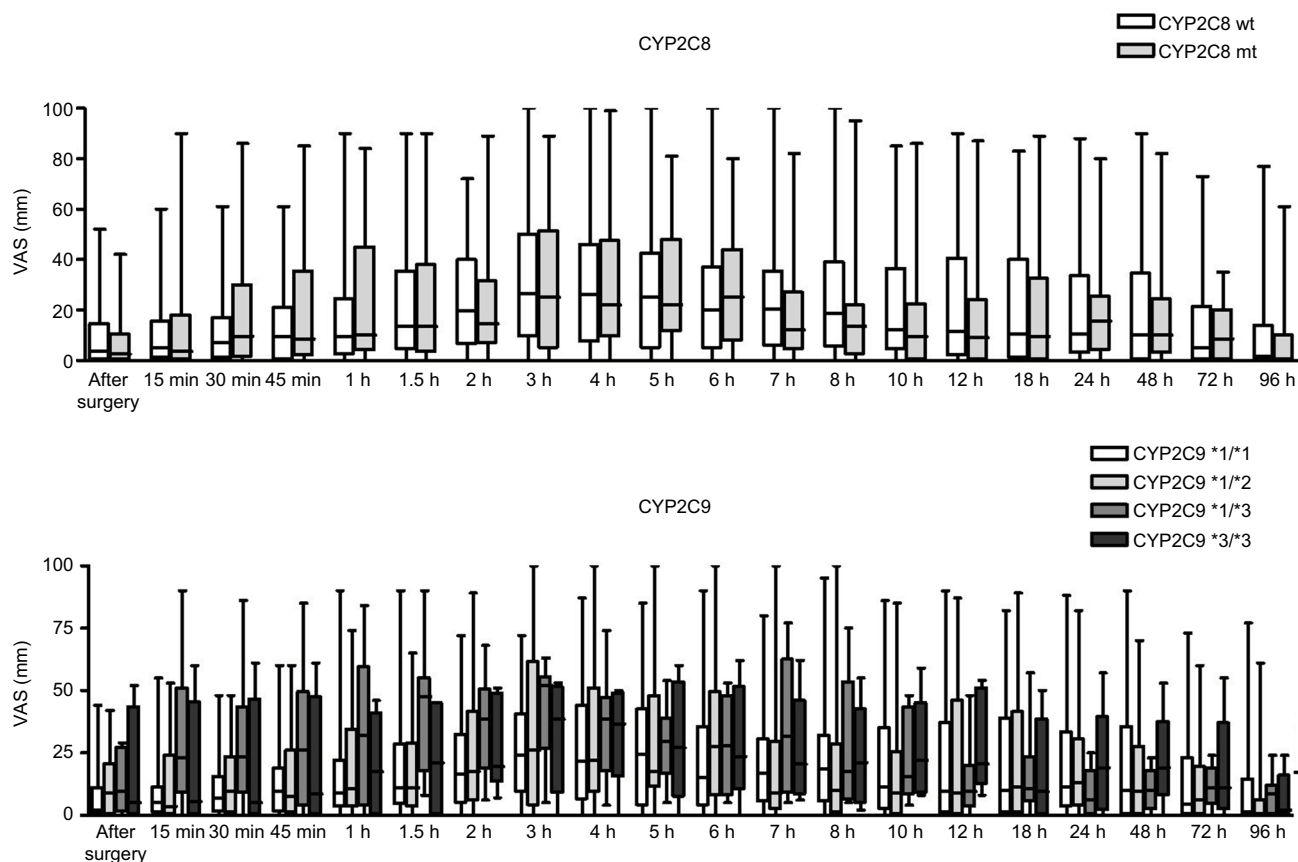


Figure 2 Postoperative pain scores reported by volunteers.

Notes: VAS of self-reported postoperative pain scores after lower third molar surgeries assessed at 0.25, 0.5, 0.75, 1, 1.5, 2, 3, 4, 5, 6, 7, 8, 10, 12, 18, 24, 48, 72 and 96 hours. Scores could range from 0 to 100 mm, with larger scores indicating increased pain. The top frame refers to the *CYP2C8**3- and the lower frame to *CYP2C9*-bearing volunteers.

Abbreviations: CYP, cytochrome P450; VAS, visual analog scale; mt, mutant; wt, Wild type.

but not the clinical control of inflammatory signs after surgical procedures. Some articles have shown pharmacogenetic impacts on large surgeries, such as the need for prior genetic analysis of patients before bariatric surgeries, regarding the lower use of opioids and lower rates of postoperative pain;²⁹ The study of Murto et al³⁰ shows the impact of *CYP2C9**3 on lower postoperative pain rates after adenotonsillectomy surgeries in children taking celecoxib.

It was speculated that postoperative pain would be effectively managed by piroxicam (20 mg daily for 4 days) for all genotypes, but that all volunteers with *CYP2C8**3 mt and *CYP2C9* genotypes might have experienced less postoperative pain. It was also possible that adverse side effects would be decreased in volunteers with the wt alleles when compared to volunteers with the mutant alleles. In other words, individuals with mutant alleles for *CYP2C8**3

and *CYP2C9* might have less pain or more side effects when compared to individuals with wt alleles for *CYP2C8*3* and *CYP2C9*. However, postoperative pain scores among the volunteers were not different when comparing the groups of both *CYP2C8*3* and *CYP2C9*.

Jensen et al³¹ noted that patients undergoing postoperative pain reporting <44 mm (VAS) tend to describe their pain as mild or having minimal impact on their daily activities. In this study, volunteers reported pain scores on average <40 mm. Similarly, Rollason et al¹ reported that the average postoperative pain score reported by their volunteers was <44 mm, demonstrating effective postoperative pain management that was independent of their *CYP* genotype. These data are clinically important since, for this type of surgery, with signs of acute pain and punctal inflammation that tend to cease within 2–3 days, patients presenting the ancestral version of *CYP2C8*3* and *CYP2C9* do not require piroxicam dose adjustment.

The main importance of pharmacogenetic clinical studies after the consumption of NSAIDs is the occurrence of gene-related changes in pharmacokinetics producing side effects rather than a change in the efficacy profile. In general, carriers of mutations in genes encoding drug-metabolizing enzymes, when treated with the usual doses of a drug metabolized by the affected enzyme, tend to have increased plasma levels of the drug, faster rate of absorption relative to its elimination and increased frequency and severity of adverse reactions with the use of these drugs.^{4–7,13} Adverse side effects noted by the surgeon or the volunteers were rare in this study, with only two individuals reporting mild adverse reactions related to consuming their daily 20 mg dose of piroxicam. Thus, no correlations were found between genotype and adverse side effects in this regimen of prescription for the control of inflammatory signs after surgeries of lower third molars with piroxicam.

Other studies with large samples report increased risks of gastrointestinal bleeding when patients consuming NSAIDs carried the *CYP2C9*2* and *CYP2C9*3* mutant alleles,^{1,5,32} although they state that this risk of gastrointestinal bleeding is clearly dose related and the pharmacokinetics shows greater exposure to these drugs in carriers of the *CYP2C9*2* and **3* alleles. Recommendations for dose adjustments in those patients are currently only for celecoxib and flurbiprofen.¹ Martinez et al genotyped *CYP2C9* in subjects who consumed various NSAIDs that are metabolized by *CYP2C9* (ibuprofen, indomethacin, celecoxib, valdecoxib, lornoxicam, tenoxicam, meloxicam and piroxicam) and other NSAIDs not considered *CYP2C9* substrates (salicylate and acetaminophen).⁵ These

authors concluded that the association of the variant *CYP2C9* alleles with the risk of gastrointestinal bleeding showed a gene–dose effect and was increased in patients receiving drugs metabolized mainly by *CYP2C9* when compared to the wt allele, suggesting that genotyping *CYP2C9* might identify subgroups of individuals potentially at risk of acute gastrointestinal bleeding.^{14,15,32,33} In the current study, only the patients' reports were used as a parameter of gastrointestinal discomfort; no endoscopic examination was performed to investigate the asymptomatic gastrointestinal side effects, and thus, small lesions may have occurred that did not affect the daily routine of the patients.

Otherwise, within the postoperative parameters examined, the volunteers' alleles for *CYP2C8*3* or *CYP2C9* were not correlated with any variation in the anti-inflammatory effects of piroxicam, indicating that piroxicam is capable of controlling inflammation regardless of their genotype. Other uninvestigated factors can interfere with inflammation such as postoperative dental hygiene around the extraction site.

Volunteers who underwent lower third molar surgeries, regardless of their *CYP* genotype, achieved satisfactory relief from postoperative pain and inflammation, with minimal risk of adverse side effects. Therefore, even for slow metabolizers of piroxicam, a daily 20 mg oral dose of piroxicam can be effective while minimizing adverse side effects after invasive oral surgery.

The impact of genetic variants on drug responses to control inflammatory signs and adverse effects of NSAIDs is clear since giving the “right drug” to the patient is a major goal in the era of medicine excellence. Further investigations with other NSAIDs used in the postoperative period of lower third molar surgeries are necessary because this group of drugs presents considerable variations regarding the genetic influence of *CYP*.

Conclusion

In general, slow metabolizers of piroxicam, namely, volunteers with mutant alleles, were indifferent from normal metabolizers with the wild-type alleles and therefore did not require specialized piroxicam doses to manage postoperative pain and inflammation after surgical removal of the lower third molar.

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Disclosure

The authors report no conflicts of interest in this work.

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