



Short Communication

## Prevalence of codon 72 P53 polymorphism in Brazilian women with cervix cancer

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### Abstract

The p53 codon 72 polymorphism seems to be associated with HPV-carcinogenesis, although controversial data have been reported. A series of Brazilian women with cervix carcinomas were analyzed. Ninety-nine (67%) of 148 women were found to be homozygous (arg/arg) for the arginine polymorphism, and 49 (33%) were heterozygous (arg/pro). This polymorphism may be an important determinant of the risk for cervix cancer, but does not seem to be sufficient for carcinogenesis.

*Key words:* p53 polymorphism, codon 72, genetic polymorphism, cervix cancer, human papillomavirus.

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Cancer of the uterine cervix is the second most frequent malignancy in women in developing countries, and responsible for substantial morbidity and mortality worldwide. Almost all cervical carcinomas harbor HPV-DNA sequences, and generally the viral E6 and E7 oncoproteins are expressed in these carcinomas (Walboomers *et al.*, 1999). The E6 and E7 of the oncogenic HPV types bind to p53 and pRb proteins, respectively. The fact that the E6 protein from the high-risk HPV can induce degradation of p53 has led to the proposal that this p53 inactivation pathway could play a key role in cervical carcinogenesis (Syrjänen and Syrjänen, 1999; Brenna and Syrjänen, 2003).

Since its detection in 1987, several studies have linked genetic polymorphism of the p53 codon 72 to carcinogenesis and progression of cervical cancer. A person can carry one of two variations of the p53 gene in codon 72: either p53 arg or p53 pro. It was suggested that oncoprotein E6 inactivates p53arg(72) more easily than pro(72), bearing some association with the outcome of HPV infections. Indeed, it has been proposed that women

who are homozygous for p53arg might have a higher susceptibility to the effects of oncogenic HPV types (Storey *et al.*, 1998; Maciag and Villa, 1999, Makni *et al.*, 2000).

The material of the present study consists of 148 Brazilian women diagnosed and treated for HPV-positive invasive cervix cancer (clinical stages I-III) in two clinics: a) Maternity Hospital Leonor Mendes de Barros of the São Paulo State Health Secretariat; and b) Women's Health Care Center of the State University of Campinas. The Ethics Committees of both hospitals approved the research plan, and written informed consent was obtained from all patients. Diagnostic biopsies from all women were available for the study. The samples were fixed in formalin, embedded in paraffin, and processed for 4- $\mu$ m-thick HE-stained sections. The histological diagnosis of carcinoma was confirmed according to routine procedures.

The demonstration of a codon 72 p53 genetic polymorphism was performed in DNA extracted from paraffin-embedded sections, using the Polymerase Chain Reaction (PCR) technique. In brief, 4- $\mu$ m sections of the tumor were treated with xylene, ethanol, proteinase K, phenol and chloroform. Pellets were resuspended in milliQ water and stored at -20 °C. Then, 5  $\mu$ L aliquots of DNA were sub-

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jected to PCR, using Master Mix (Eppendorf, Düsseldorf, Germany) together with the following primers:

ArgF: 5'-TCC CCC TTG CCG TCC CAA -3'

ArgR: 5'-CTG GTG CAG GGG CCA CGC-3' (144

bp)

ProF: 5'-GCC AGA GGC TGC TCC CCC-3'

ProR 5'-CGT GCA AGT CAC AGA CTT-3' (177bp)

$\beta$  globin F: 5'-CAA CTT CAT CCA CGT TCA CC-3'

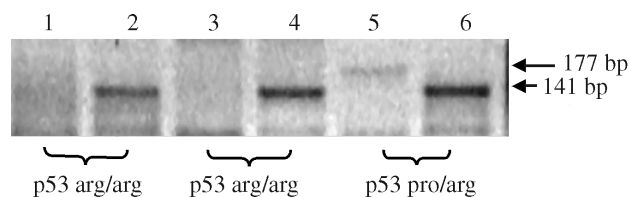
$\beta$  globin R: 5'-GAA GAG CCA AGG ACA GGT AC-3'

PCR was performed in separate tubes for arg, pro alleles, and  $\beta$  globin; bands were separated in 2% low-melting point agarose stained with ethidium bromide and analyzed using the Kodak EDAS 120 software in order to increase sensitivity. All samples were previously analyzed for  $\beta$  globin amplification, in order to check the quality and quantity of DNA. Thermal cycling was accomplished by using a GenAmp PCR System 9700 (Applied Biosystems). The cycling profile, as well as the agarose gel electrophoresis, followed exactly the protocol used by Soultziz *et al.* (2000) (Figure 1).

Unconditional logistic regression analysis was used to calculate odds ratios (ORs) as measures of association between p53 polymorphism and the clinical stage, with respective 95% confidence intervals (CIs). Confounding control was done by multiple logistic regression models, resulting in adjusted ORs. The statistical analyses were made with the SAS version 8.2 software.

The histological diagnosis of squamous cell carcinoma (SCC) was confirmed in 144 (97%) and of adenocarcinoma in 4 (3%) of the cases. Distribution according to clinical stage was as follows: 40 (27%) stage I, 49 (33%) stage II, and 59 (40%) stage III. Among the 148 women, 99 (67%) were homozygous for the arginine polymorphism (arg/arg), 49 (33%) were heterozygous (arg/pro) and none was homozygous for (pro/pro).

Table 1 shows the distribution of the p53 polymorphism according to the clinical stage. There was no significant difference in the distribution of the polymorphism between the clinical stages. Multiple logistic models containing also the interaction terms evaluated the joint association of polymorphisms with clinical stages, but this analysis did not show evidence suggesting a relationship between the variables and p53 polymorphism.



**Figure 1** - PCR amplification of the p53 codon 72 arg allele (144 bp) and pro allele (177 bp). Agarose and ethidium bromide, analyzed by Kodak EDAS 120.

**Table 1** - Distribution of p53 polymorphism in Brazilian women according to the FIGO stages.

Stages	p53 polymorphism				crude OR (95% CI)	adjusted* OR (95% CI)
	arg/arg		arg/pro			
	n	(%)	n	(%)		
I	27	(27)	13	(26)	Ref	Ref
II	32	(33)	17	(35)	0.9 (0.3-2.2)	0.9 (0.3-2.1)
III	40	(40)	19	(39)	1.0 (0.4-2.4)	0.9 (0.3-2.1)
Total	99	(67)	49	(33)		

\*Adjusted for the other variables in the Table.

Brazil is a multiracial tropical developing country with high incidence and mortality rates of cervix cancer, 18.32 and 4.58 per 100.000, respectively (Brasil, 2003). In the general population, a polymorphism in p53 occurs as a result of a proline to arginine transition at codon 72 of exon 4. There is no accurate information about the frequency of that polymorphism in normal Brazilian women, neither in women with cervix cancer. However, Brazilian authors have previously evaluated 82 cases of oral squamous cell carcinoma and 82 controls with similar population characteristics, but with a different tumor biology. They found a genotype distribution of 37.8% (31) arg/arg, 54.8% (45) arg/pro, and 7.4% (6) pro/pro in the tumor group, and of 40.2% (33), 54.8% (45), and 4.9% (4), respectively, in the normal population (Drummond *et al.*, 2002).

It can be speculated that the genetic background of the host influences the persistence of HPV infection, which could mediate the genetic susceptibility for cervix cancer. Storey *et al.* (1998) presented data suggesting that individuals who are homozygous arginine display a higher susceptibility of p53 to degradation by the high-risk HPV E6 *in vitro*. Furthermore, they found that individuals homozygous for arginine were about seven times more susceptible to HPV-associated cervix carcinoma than heterozygotes.

However, the published data on the prevalence of p53 polymorphism in cervix cancer patients are controversial. As anticipated, the ethnic group characteristics seem to be an important reason for discrepancies in the frequency of this polymorphism. Thus, the frequency of the arginine allele increases with latitude, while the proline allele shows the opposite effect (Beckman *et al.*, 1994).

In Asian women, the frequency of p53arg was found to be relatively low: 42% (Baek *et al.*, 2000) and 40% (Kim *et al.*, 2001) in Korean women; 33% (Yamashita *et al.*, 1999) and 44.8% (Nishikawa *et al.*, 2000) in Japanese women; 31% in Hong Kong women (Wong *et al.*, 2000); and 27% in Indian women (Bhattacharya *et al.*, 2002). Similar results were reported in Southern Europe. In Israel, the frequency was 34.8% in Jewish women, 30.3% in North African women, and 10.8% in those of other origins (Arbel-Alon *et al.*, 2002). The results of two series of obser-

vations on Italian women were controversial: 34% (Tenti *et al.*, 2000) and 76.7% (Zehbe *et al.*, 2001). A number of studies on ethnically diverse groups of patients with cervix cancer from Northern Europe have revealed a higher frequency of p53arg than that observed in Asian and South European women. Thus, the frequency of p53arg observed in white English women was 54% (Rosenthal *et al.*, 1998) and 69.4% (Brady *et al.*, 1999), in Dutch women (from the Netherlands) 62% (van Duin *et al.*, 2000), and in Swedish women 63.9% (Zehbe *et al.*, 2001).

Similar reports are available on cervical cancer patients from America, suggesting that in South and North America the frequency of p53arg is higher than in Central America. The results of a study carried out in Peruvian women revealed a 50.4% frequency of p53arg (Klug *et al.*, 2001), and in two series of American women, the figures were 55.9% (Madeleine *et al.*, 2000) and 67% (Malcolm *et al.*, 2000), respectively. In Central America, results are available from Mexico, showing a 45% frequency of p53arg (Suarez-Rincon *et al.*, 2002). Against this background, the 67% prevalence of the arg/arg polymorphism in Brazilian women seems to be high, but it is still consistent with the American data reported (Madeleine *et al.*, 2000; Malcolm *et al.*, 2000; Klug *et al.*, 2001; Suarez-Rincon *et al.*, 2002).

The controversial data found by the different studies have been attributed to ethnic differences, but other potential confounding factors should be considered as well. These could include the sample size, the source of DNA, and the detection techniques used. Another important reason for these discrepant results could be misclassification of the p53 polymorphism, due to inter-laboratory variations in protocols, affecting the ability to detect p53 polymorphisms. When the p53 polymorphism was analyzed in three different laboratories, the differences in arg/arg genotype compared to other forms were not significant. After exclusion of the discordant genotype, however, the association of arg/arg p53 with cervix cancer was significant, with an OR = 8.0 (95% CI .3-28.5) (Makni *et al.*, 2000).

In conclusion, such variations reflect the geographic spread of HPVs. The p53 polymorphism may be one of the important determinants of the risk for cervix carcinoma. However, it does not seem to be sufficient to induce cervix carcinogenesis or to determine the progression of the disease. The role of the genetic susceptibility to HPV infections and cervix cancer merits further investigation.

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