

# Analysis of Polymorphism rs1042522 in *TP53* Gene in the Mothers of Twins and of Singletons: A Population-Based Study in Rio Grande do Sul, Brazil

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Although the birth of twins has always attracted attention, there are no known genetic or environmental factors that can determine the birth of monozygotic (MZ) twins. And even for dizygotic (DZ) twins, genetic influences are not completely understood. A previous study from our group has shown that the C allele of polymorphism rs1042522 in the *TP53* gene was more frequent in the mothers of twins than in the mothers of singletons in a small village in South Brazil. In order to clarify whether this was an isolated factor, we performed a population-based, observational case-control study. Samples were selected from a state-funded program of paternity investigation. Samples were considered cases when two of the children had the same date of birth, whereas controls were those samples in which at least two children were born in different dates. The first subsequent sample fulfilling control criteria was included after each case. From 2007 to 2013, 32,661 records were searched and 283 (0.9%) twins were found (119 MZ and 164 DZ). Genotypic and allele frequencies were not different between mothers of twins or mothers of singletons. However, mothers of MZ twins showed a higher frequency of GG genotype and lower frequency of the C allele when compared to mothers of DZ twins. Also, the proportion of MZ twins (42%) was higher than usually reported (30%). Finally, the proportion of twins found in this study seems to be more realistic, as this sample was allegedly not from users of assisted reproduction techniques.

■ **Keywords:** twins, mutation, rs1042522, *TP53*

Although the birth of twins has always attracted attention, there are no known genetic or environmental factors that can determine the birth of monozygotic (MZ) twins. The existence of hereditary factors is suggested by familial aggregation, especially for dizygotic (DZ) twins. Although the genetic influences are not completely understood, recent evidence points to the involvement of genetic variants involved in gonadal response to FSH, such as *FSHB* and *SMAD3* (Mbarek et al., 2016). However, the gene for DZ twinning has yet to be mapped in humans, and even the mode of inheritance is not known (Hoekstra et al., 2008). MZ twins, on the other hand, seem to happen by chance only, without influence of ethnicity, maternal age, parity, or nutritional state (Tong & Short, 1998).

An increase in the incidence of DZ twins has been observed worldwide, especially due to the use of assisted reproduction and a residual effect of the prolonged use of contraceptives (Hall, 1996). In the United States, the twinning rate was stable at 32 per 1,000 live births in 2006 (Chauhan et al., 2010), although more recent data suggested figures

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of 16.9 per 1,000 in 2011 (Martin et al., 2012). In Europe, the twinning rate has wide variation across countries, ranging from 16.1 per 1,000 in England and Wales to 21.2 per 1,000 in France (Pison et al., 2015). In Brazil, data from 1995 to 1998 showed rates of 19.51 per 1,000 births for DZ twins, whereas the MZ twinning rate was of 4.5 per 1,000. This latter number is similar to the 4.1 per 1,000 previously described in the country (Colletto et al., 2001), and to figures reported elsewhere in the world (Macfarlane & Blondel, 2003).

In Rio Grande do Sul (RS), the southernmost state of Brazil, there is a city called Candido Godói whose population presents a high rate of twinning, estimated at 15 per 1,000 births. Epidemiological data suggest a founder effect, dating from the late 19th century. Genetic analysis of *TP53* and its regulators (*MDM2*, *MDM4*, and *HAUSP*) showed an association of polymorphism rs1042522 in *TP53* and twinning in this population (Tagliani-Ribeiro et al., 2012). An additional study on twinning and *TP53* was performed by Huang et al. (2015) in African Americans, but the association they found was with a different polymorphism (rs8079544). So far, there is no data on the association of polymorphism rs1042522 in *TP53* and twinning in other regions, and it is debatable whether this is an isolated factor.

The Laboratory of Paternity Investigation (LPI) is a state-based laboratory in RS that receives samples from low-income individuals for paternity investigation paid by the public defense system. On average, 400 cases from all over the state are analyzed per month. Considering that this population-based sample is presumably free of influences caused by fertility treatments, we decided to study the frequency of polymorphism rs1042522 in *TP53* gene, and its relation to the birth of twins.

## Material and Methods

This was a population-based, observational case-control study. Samples were selected from the LPI database of the period 2007–2013 from those where there was no doubt about maternity. Out of the 32,661 records, 583 were selected to fulfill criteria as cases or controls. Samples were considered cases when two of the children had the same date of birth, whereas controls were those samples in which at least two children were born in different dates. The first sample fulfilling control criteria was included after each case. For each sample, the following data was collected: mother's birth date and city of origin, and children's birth date.

DNA was extracted from archived FTA Cards (Whatman) using a commercial solution, according to the manufacturer's instruction (FTA Solution, Whatman). The presence of polymorphism rs1042522 in *TP53* gene was determined in DNA samples from the mothers using TaqMan Assay C\_2403545\_10, as described by Tagliani-Ribeiro

et al. (2012). Zygosity of same-sex twins was determined from data used in the panel of genetic markers for paternity investigation (CSF1PO, D13S317, D16S539, D18S51, D19S433, D21S11, D2S1338, D3S1358, D5S818, D7S820, D8S1179, FGA, TH01, TPOX, VWA, and amelogenin).

Statistical analysis using a chi-square test was performed in SPSS version 18.0,  $p < .05$  considered significant. Approval from the local ethics committee was obtained without the need for informed consent as all data were treated anonymously.

## Results

From January 2007 to December 2013, LPI performed 32,661 paternity investigations. Samples originated from 12 cities that were grouped into five regions (Figure 1). Out of these, 283 (0.9%) were from mothers of twins. Therefore, the other 300 samples were included as controls (mothers of singletons). The mean age of mothers of twins at the time of twins' birth was 26.4 years (MZ = 25.4 and DZ = 27.07), whereas the mean age of mothers of singletons at the time of the first child's birth was 24.5 years.

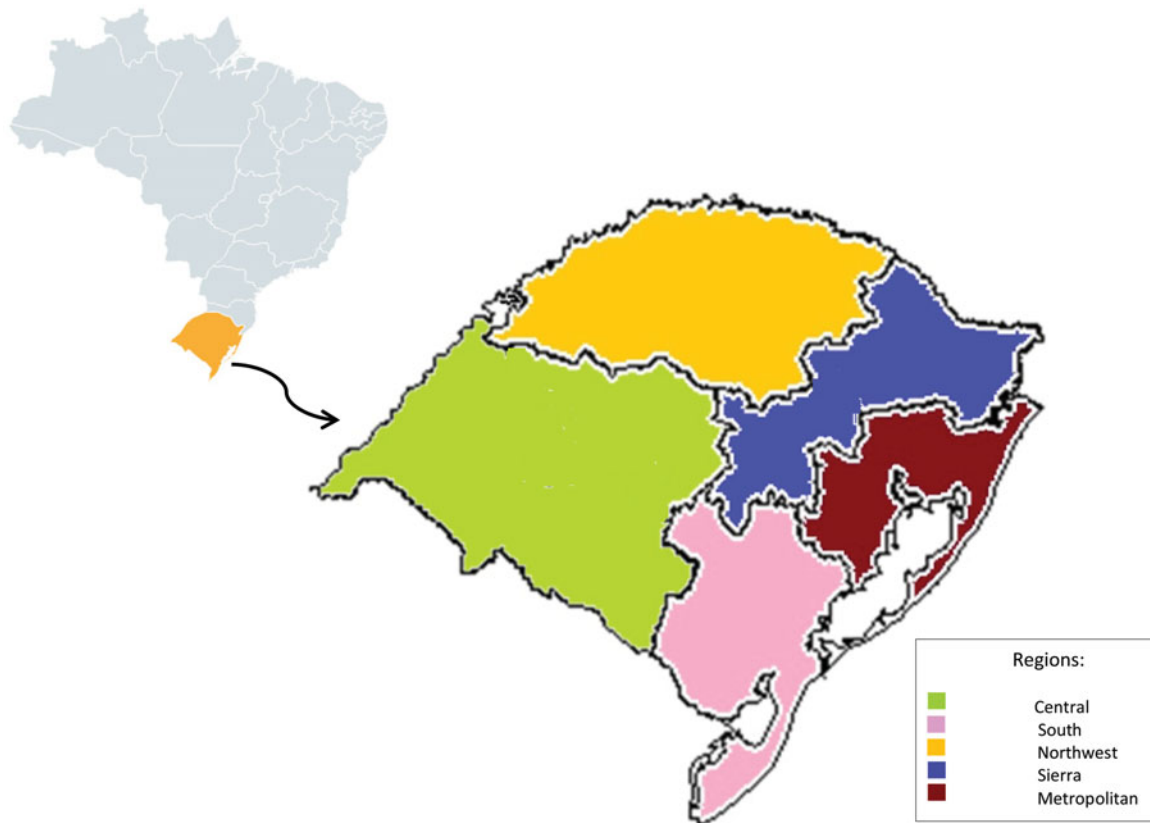
Among cases, the percentage of MZ twins was 42%, differing from the estimated one-third referred to in the literature ( $p = .04$ ). The distribution of twin births as the percentage of MZ twins per region can be observed in Table 1. There was no statistically significant difference across regions regarding the percentage of twins ( $p = .744$ ). However, the Sierra and Northwest regions showed a smaller percentage of MZ twins compared to the other regions ( $p = .0004$ ), although with values closer to those described in the literature (30%).

Data analysis of polymorphism rs1042522 in *TP53* gene in the mothers of twins and of singletons did not show a statistically significant difference in genotypic (Table 2,  $p = .371$ ) or allelic frequencies (Table 2,  $p = .187$ ).

However, when analyzed by zygosity, there was a difference in genotypic and allelic frequencies between mothers of DZ and MZ twins (Table 3). Genotype GG was more frequent than expected in the mothers of MZ twins ( $p = .028$ ), whereas the C allele was diminished ( $p = .016$ ) in this same group of individuals. The odds ratio for allele G was 1.61 and for allele C it was 0.62 (SE 0.26).

## Discussion

Although there is evidence of familial susceptibility for twinning, the genetic factors involved are unknown. Among studied candidate genes are growth differentiation factor 9 (*GDF9*) and bone morphogenetic protein 15 (*BMP15*), essential for normal human fertility and with a prominent role in follicle growth and ovulation (Moore et al., 2004). Recently, a genome-wide association study identified genetic risk variants for DZ twinning on a region near *FSHB* and *SMAD3*. These genes are involved in gonadal responsiveness to FSH and might contribute

**FIGURE 1**

(Colour online) Regions of sample origin in Rio Grande do Sul (RS). The inset shows the location of RS in Brazil.

**TABLE 1**

Distribution of Twins (Both MZ and DZ) and of MZ Twins in Different Analyzed Regions

Region	Cases (MZ + DZ twins) n (%)	MZ twins n (%)
Metropolitan	124 (43.8)	54 (45.4)
Central	33 (11.7)	18 (15.1)
Sierra	32 (11.3)	8 (6.7)*
Northwest	60 (21.2)	21 (17.6)*
South	34 (12.0)	18 (15.1)
Total	283 (48)	119 (42)

Note: \* $p = .0004$ .

**TABLE 2**

Distribution of Genotypic and Allelic Frequencies of Polymorphism rs1042522 in *TP53* Gene in the Mothers of Twins ( $n = 285$ ) and of singletons ( $n = 300$ )

	Mothers of twins n (%)	Mothers of singletons n (%)
Genotype		
GG	140 (49.5)	131 (44.0)
GC	116 (41.0)	136 (45.0)
CC	27 (9.5)	33 (11.0)
Allele		
G	396 (69.80)	398 (66.3)
C	172 (30.20)	202 (33.7)

to crucial aspects of reproductive capacity (Mbarek et al., 2016).

A previous study from our group has shown that the C allele of polymorphism rs1042522 in *TP53* gene was more frequent in the mothers of twins than in the mothers of singletons in Candido Godoi, a small village in South Brazil (Tagliani-Ribeiro et al., 2012). However, the present study, analyzing a population-based sample, did not show a difference in allelic or genotypic frequencies between the mothers of twins and singletons. Nevertheless, a higher frequency of the C allele was found in the mothers of MZ twins as compared to the mothers of DZ twins. It is therefore, possible that the previous difference was in fact

due to an excess of MZ twins' mothers in that sample. Unfortunately, zygosity was not performed by molecular studies in the previous work. Yet, there is the possibility that the data found in Candido Godoi by Tagliani-Ribeiro et al. (2012) reflected a founder effect that cannot be extended to other populations. Unfortunately, no data on ethnicity of our sample is available to tease out whether population stratification could be the basis of this difference. Nevertheless, data on genetic admixture of Latin American populations (Ruiz-Linares et al., 2014) showed a marked genetic homogeneity in the southernmost part of Brazil — the same origin of samples for the present study. In

**TABLE 3**  
**Distribution of Genotypic and Allelic Frequencies of Polymorphism rs1042522 in TP53 Gene in Mothers of MZ (n = 119) and DZ (n = 166) Twins**

	Mothers of MZ twins n (%)	Mothers of DZ twins n (%)
Genotype		
GG	70 (59.0)	70 (43.0)
GC	40 (33.5)	76 (46.0)
CC	9 (7.5)	18 (11.0)
Allele		
G	180 (75.2)	216 (65.8)
C	58 (24.8)	112 (34.2)

addition, in the present work we could not exclude the presence of twins in the families of singleton mothers, as no data regarding other relatives were investigated (even though the previous birth of twins was ruled out).

Despite the lack of information in some relevant aspects, we chose to study this particular population as it can be considered free of influences caused by fertility treatments, since these are women using a public service for paternity investigation, limited to low income users, who do not have access to private fertility treatments. In our state, there is only one public service for assisted reproduction, and entrance to this program is difficult and limited to stable couples after assessment by a social worker. Therefore, we can assume that at least in its vast majority, this sample is free from influence of fertility treatments.

To the best of our knowledge, this is the second study to investigate the role of *TP53* in twinning. Despite being extensively known for its role as a tumor suppressor (Levine et al., 2006), the *TP53* gene also plays an important role in fertility. Forms of the p53 family (including p63 and p73) are involved in the maintenance of follicle pool, genomic integrity of germ cells, and blastocyst implantation (Hu et al., 2007). One of *TP53*'s most studied variants is polymorphism rs1042522, which changes a cytosine into a guanine, thus replacing a proline with an arginine in residue 72 of the protein (p.P72R).

Studies in non-human primates suggest that the C allele (p.P72) is ancestral, although the G allele (p.R72) can reach frequencies over 0.5% in certain populations (Whibley et al., 2009). This change results in mild functional differences in the protein, and the variant p.P72 (C allele) is more efficient in initiating senescence and arresting the cell cycle, whereas the p.R72 (G allele) variant is more active in apoptosis induction and in suppressing cell transformation (Dumont et al., 2003). Kang et al. (2009) investigated the impact of such changes in the rates of embryo implant and successful pregnancy after in vitro fertilization. In young patients, p.P72 seems to be a risk factor for implantation failure. These data corroborate our study, as the C allele was under-represented in mothers of MZ twins.

Maternal age is a factor usually associated with twinning, although not with MZ twins (Aston et al., 2008). A positive

association was described for DZ twins and maternal age by several groups (Lambalk et al., 1998). In the present study, no influence of maternal age was observed, even when comparing MZ and DZ mothers. This may be due to the sample type used in this study, formed mainly by young women having unplanned pregnancies. Therefore, a bias may be introduced, especially regarding DZ twins.

Also on this aspect, it is worth noticing the apparent excess of MZ twins in our sample. The reasons for this are not completely understood, because the risk factors for MZ twinning are not clear. Familial aggregation of MZ twins is rare and its cause may be due to a single gene related to the integrity of zona pellucida or the maintenance of cell-cell connection (Hall, 1996). It is assumed that MZ twins comprise 30% of all twin births (Aston et al., 2008), but in the present sample they accounted for almost 42% of cases (119/283).

Finally, the number of twin births found in this population-based sample (283) corresponds to only 0.9% of all paternity investigations analyzed at the LPI. Since the twinning rate for the state of RS is estimated to be 2% (DATASUS, 2015), one would expect that among the about 32,661 paternity investigations, 653 were of twins. However, the number found was almost half this value. Since there is no reason to suppose that mothers of twins seek state-paid paternity investigation in a different proportion compared to mothers of singletons, it may be inferred that the rate found in this study is more representative of the real twinning rate for RS, without the impact of assisted reproduction techniques. This hypothesis could be evaluated if data on the frequency of assisted reproduction among parents of twins were known or, if historical records of twin births were available for the state. Unfortunately, none of these figures are publicly available in Brazil. The oldest record of twin births available at DATASUS dates back to 1994 and indicates a proportion of 1.77% twin births in RS.

In brief, the present study investigated the frequency of polymorphism rs1042522 in *TP53* gene in mothers of twins and singletons using a population-based sample of women seeking state-funded paternity investigations. Genotypic and allele frequencies were not different between groups, but mothers of MZ twins showed a higher frequency of GG genotype and lower frequency of the C allele when compared to mothers of DZ twins. Also, the proportion of MZ twins seems to be elevated in this sample and may represent a more realistic figure on twinning rates.

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