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Maternal MTHFR polymorphisms and risk of spontaneous abortion

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Abstract
Objective. To assess the association between intake of folate and B vitamins and the incidence of spontaneous abortion (SA) according to the maternal methylenetetrahydrofolate reductase (MTHFR) polymorphisms (677 C>T and 1298 A>C). Material and Methods. We conducted a nested case-control study within a perinatal cohort of women recruited in the state of Morelos, Mexico. Twenty-three women with SA were compared to 74 women whose pregnancy survived beyond week 20th. Intake of folate and B vitamins respectively, was estimated using a validated food frequency questionnaire. Maternal MTHFR polymorphisms were determined by PCR-RFLP and serum homocysteine levels by HPLC. Results. Carriers of MTHFR 677TT and 1298AC genotypes respectively showed an increased risk of SA (OR 677TT vs. CC/CT=5.0; 95% CI: 1.2, 20.9 and OR 1298 AC vs. AA=5.5; 95% CI: 1.1, 26.6). Conclusions. Our results support the role of MTHFR polymorphisms as a risk factor for SA, regardless of dietary intake of B vitamins.

Key words: MTHFR; 677 C>T and 1298 A>C polymorphisms; spontaneous abortion; Mexico

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Spontaneous abortion (SA) is defined as the loss of a fetal product before 20 weeks of gestation. Lifestyle, diet, and, more recently, maternal genetic characteristics have been proposed as determinants of SA. Specifically, maternal and paternal smoking and maternal alcohol and coffee consumption during pregnancy have been associated with a higher risk of SA. In addition, low folate levels in maternal serum and suboptimal folate metabolism, indicated by the presence of hyperhomocysteinemia, have also been associated with recurrent abortions. The relationship of SA with intakes of other B vitamins that are involved in folate metabolism, i.e., B2, B6 and B12, has not been well established.

Methylenetetrahydrofolate reductase (MTHFR), a key folate-metabolizing gene, is located on chromosome 1. Several non-synonymous single nucleotide polymorphisms (SNPs) are present in the coding region of the gene including positions 677 and 1298. The most studied SNP is the 677 C>T substitution that results in an amino acid change from alanine to valine at codon 225. The MTHFR 1298A>C polymorphism results in a substitution of glutamate with alanine.

MTHFR enzyme activity is reduced by 35% among the 677CT carriers and by 50% to 70% among 677TT carriers. The function of polymorphism 1298A>C has not been consistently demonstrated; however carriers of the heterozygous genotypes (677CT/1298AC) show MTHFR activity similar to that in MTHFR677TT carriers. The reduction in MTHFR activity increases homocysteine levels and in turn reduces the availability of the DNA methyl groups.

The associations reported to date between maternal MTHFR 677 C>T and 1298 A>C polymorphisms and risk of SA have been inconsistent. Some authors reported as much as a three-fold increase in SA risk among women carriers of the 677CT genotype compared to women with the 677CC or 677CT genotypes, while other authors found no association at all. An increased risk for SA was also observed in fetal carriers of the MTHFR 677T allele.

The prevalence of the MTHFR 677TT genotype is higher among Mexicans (36-48%) compared to Europeans (7.8-18%). In contrast, a much lower frequency of the MTHFR 1298C allele has been observed in the Mexican population (0 to 2.4% vs. 7.5 to 11.5%) than in Europeans.

The objective of this study was to evaluate the risk of SA according to maternal MTHFR polymorphisms (677 C>T and 1298 A>C) and other lifestyle risk factors in a cohort of Mexican women residing in the state of Morelos in Mexico.

Material and Methods

We conducted a nested case-control study within an ongoing perinatal cohort of women that began in January 2001; the cohort had been formed to evaluate the effect of environmental exposure to organochlorine on infant neurodevelopment. Study participants were recruited from four municipalities in the state of Morelos, Mexico, during premarital counseling mandated by Mexican law. To be eligible for the study, women had to be non-lactating and without a history of chronic diseases. A detailed description of the methodology and cohort follow-up is reported elsewhere.

Women were contacted every six weeks by telephone or household visit and asked to report a ten-day delay in their menstrual period to confirm a pregnancy status with a pregnancy test, or clinically document the presence of a SA.

Twenty-three women with SA (pregnancy that ended spontaneously before 20 weeks) with a median gestational age of 10 weeks (min=6 and max=17), were identified among a group of 456 women who became pregnant before July 2004, and 74 women whose pregnancies lasted beyond 20 gestational weeks, with a median gestational age of 37.2 (min=21 and max=42), were randomly selected as controls.

Home interviews were conducted on both women and their husbands/partners to elicit sociodemographic information (e.g., partners’/husband’s occupation), reproductive history, and dietary habits before pregnancy, alcohol consumption prior to conception and tobacco use. During the baseline evaluation, each woman’s weight and height were measured to calculate her body mass index (BMI), and a blood sample (7 ml) was obtained to determine her MTHFR genotypes and homocysteine serum levels.

This study was approved by the Institutional Review Board of the National Institute of Public Health of Mexico. Informed consent was obtained from study participants and their husbands.

Genotyping of the MTHFR 677 C>T and 1298 A>C Polymorphisms

Genomic DNA was extracted using a QIAamp DNA Blood Mini Kit (QUIAGEN, Valencia, CA). DNA amplification and genotyping of MTHFR 677 C>T and 1298 A>C were performed by polymerase chain reaction (PCR-RFLP) according to the protocol suggested by Chen et al. and Weisberg et al. For quality control purposes, for each batch of PCR-amplified samples, one
sample containing a known genotype and one negative control were analyzed. In addition, duplicate analyses of 10% of the samples, randomly selected, were performed by a technician who was blinded to the first set of results and reached Kappa correlation coefficients of 1 and 0.954 for the MTHFR 677 C>T and MTHFR A>C 1298 genotype determinations respectively.

**Dietary intake of folate and B vitamins**

Dietary intake of folate and vitamins B_6, B_12, and B_15 was estimated by means of a validated semi-quantitative food frequency questionnaire (FFQ) containing 95 different items. For each food item, there was a defined portion, and women were asked to choose from 10 consumption frequency options, which ranged from never to six times per day. The consumption of 28 individual nutrients (including total caloric intake) was calculated using the Food Intake Analysis Software 3.0 (FIAS, Texas University, USA). A detailed description of the methods used to calculate nutritional intake has been published elsewhere.

On the FFQ the following foods were considered sources of vitamins B_6, B_12, and folate: liver, beans, lentils, green beans, peas, tomatoes, spinach, lettuce, potatoes, cauliflower, broccoli, chili peppers, beets, carrots, avocados, oranges, papayas, mangos, strawberries, bananas, tangerines, melons, plums, pears, pineapples, apples, peaches, grapes, lamb, tuna, pork and steak; while the dietary intake of vitamin B_12 was based in the consumption of liver, sardines, tuna, pork, lamb, steak, fish, chocolate, beer, and red and white wines.

**Determination of serum homocysteine levels**

Total serum homocysteine levels were determined in duplicated analysis using a high performance liquid chromatography (HPLC) method previously described by Gilfix et al. Fluorescent intensities were measured with excitation and emission wavelengths set at 385 and 515 nm, respectively. Quality control (Liquicheck Homocysteine Control- Bilevel, Biorad Inc) showed 98.4% ± 2.7% and 99.1 ± 1.2%, recovery for the low and the high level, respectively. CV between duplicate sample analysis was 5.2 ±2.8%.

**Statistical analysis**

Selected maternal and paternal characteristics were compared between cases and controls. Based on the median values observed in the control group the following categories were created for: maternal age (≤ 21vs. >21 years), education (≤ 9 vs. >9 years), body mass index (≤ 22.4 vs. >22.4), and paternal age (≤ 24 vs. >24 years).

Paternal occupation was categorized as “risky” or “non-risky” according to the occupation types that have been assessed with relation to SA: poultry farmer, bricklayer, painter, maintenance worker, farmer, potter, flower-grower, pesticide applicator, and blacksmith. Due to the skewed dietary distributions, nonparametric tests were used to compare, between cases and controls, the folate and B vitamins intakes respectively. We estimated the Hardy-Weinberg equilibrium test among controls to evaluate the observed and expected frequencies of the genotypes of interest. The risk of spontaneous abortion (SA) in relation to dietary consumptions of folate and B vitamins as well as the presence of maternal methylene-tetrahydrofolate reductase (MTHFR) polymorphisms (677 C>T and 1298 A>C) was estimated using odds ratios (ORs) and their respective 95% confidence intervals (CIs) with a recessive hereditary model, and further stratify by early and late abortion (≤10 and >10 weeks). STATA (College Station, Texas, USA) statistical software version 9.0 was used to conduct all of the analyses.

**Results**

Compared with homemakers, women with paid employment had an almost three-fold borderline significant increased risk of SA (OR=2.7; 95% CI: 1.0, 7.1). Smoking (before pregnancy) as well as paternal occupation resulted in increased risks of SA respectively, however this was not significant. In contrast, maternal younger age and higher education showed non-significant protection of SA (table I).

As described in table II, non-significant higher dietary intakes of folate and B vitamins were observed among controls compared to cases, except for vitamin B_6 intake, which resulted in the border of significance (2.1 vs. 1.8 mg/d; p=0.09). Cases in contrast to controls, showed a non-significant higher median of homocysteine serum levels, as well as the MTHFR 677TT genotype (43.5% vs. 25.7%). No carriers of the MTHFR 1298CC genotype were present in this study population. The median values of homocysteine remained different (with no statistical significance) between cases and controls, after further stratification by the alleles of interest (677C: Cases=9.6 vs. Controls=10.3 µmol/L; 677T: Cases=11.0 vs. Controls=10.4 µmol/L; p=0.05. 1298A Cases=10.4 vs. Controls= 10.5 µmol/L; 1298C: Cases=9.8 vs. Controls= 8.8 µmol/L; p=0.05 ) (data not included in the table). Among the controls, neither of the genotype distributions showed any deviance from the Hardy-Weinberg equilibrium.
El estudio detectó un aumento significativo del riesgo de SA entre mujeres portadoras de los géneros MTHFR 677TT y MTHFR1298 AC, independiente de otros factores previamente asociados con un aumento de SA (uso de tabaco paterno y ocupación materna). Estos resultados apoyan los hallazgos de estudios recientes que muestran un efecto de heterozigosidad para MTHFR 677 C>T en abortos recurrentes en mujeres de edad temprana, pero deben interpretarse con precaución debido al tamaño muestral pequeño de este estudio.

El mediano de todos los vitaminas B, folatos, vitamina B12 y vitamina B12, fueron menores en mujeres con SA comparadas con los controles, aunque la diferencia no alcanzó la significación. Es importante señalar que la suplementación con multivitaminas es baja en la población mexicana,39 como fue el caso en esta pequeña población estudiada. Así, estos hallazgos deben ser replicados en estudios más grandes para evaluar la protección de estas vitaminas en el riesgo de SA.

Similar a los hallazgos en otras poblaciones mexicanas,30,31,39,40-42, se observó un alto porcentaje de nuestro estudio, con ambos casos y controles, portadores del genotipo MTHFR 1298 CC en el padre (11.4% y 9.5%, respectivamente), y una baja frecuencia del alelo 1298T (58.7% y 52.0%, respectivamente), y una baja frecuencia del alelo 1298C en la población mexicana, independientemente de otros factores ya mencionados.

La discusión anterior (OR=5.5; 95% CI:1.1, 26.6) cuando se comparan con el tipo de SA genotipo. Este resultado permaneció significativo entre abortos tardíos y no se evaluó entre abortos recurrentes debido al tamaño muestral pequeño.

**Discusión**

Este estudio detectó un aumento significativo del riesgo de SA entre mujeres portadoras de los géneros MTHFR 677TT y MTHFR1298 AC genotipos, independiente de otros factores previamente asociados con un aumento de SA (uso de tabaco paterno y ocupación materna). Estos resultados apoyan los hallazgos de estudios recientes que muestran un efecto de heterozigosidad para MTHFR 677 C>T en abortos recurrentes en mujeres de edad temprana, pero deben interpretarse con precaución debido al tamaño muestral pequeño de este estudio.

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### Table II

**DIETARY INTAKE OF FOLATE AND B VITAMINS, HOMOCYSTEINE SERUM LEVELS AND MTHFR POLYMORPHISMS.**

**Morelos, Mexico, 2001-2004**

<table>
<thead>
<tr>
<th>Variables</th>
<th>Cases (n=23)</th>
<th>Controls (n=74)</th>
<th>P values</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>P&lt;sub&gt;10&lt;/sub&gt; Median P&lt;sub&gt;90&lt;/sub&gt;</td>
<td>P&lt;sub&gt;10&lt;/sub&gt; Median P&lt;sub&gt;90&lt;/sub&gt;</td>
<td></td>
</tr>
<tr>
<td><strong>Dietary Intake</strong></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Folate, µg/d</td>
<td>131.7 341.4 514.2</td>
<td>231.6 392.9 580.8</td>
<td>0.18*</td>
</tr>
<tr>
<td>Vitamin B&lt;sub&gt;6&lt;/sub&gt;, mg/d</td>
<td>1.1 2.1 2.5</td>
<td>1.6 2.2 3.4</td>
<td>0.16*</td>
</tr>
<tr>
<td>Vitamin B&lt;sub&gt;12&lt;/sub&gt;, mg/d</td>
<td>0.9 1.8 2.7</td>
<td>1.2 2.1 3.3</td>
<td>0.09*</td>
</tr>
<tr>
<td>Homocysteine serum level (µmol/L)</td>
<td>1.9 3.2 6.3</td>
<td>1.8 3.8 8.2</td>
<td>0.53*</td>
</tr>
<tr>
<td><strong>Polymorphisms</strong></td>
<td>(%)</td>
<td>(%)</td>
<td></td>
</tr>
<tr>
<td>MTHFR 677C&gt;T</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CC</td>
<td>26.1</td>
<td>21.6</td>
<td></td>
</tr>
<tr>
<td>CT</td>
<td>30.4</td>
<td>52.7</td>
<td>0.15†</td>
</tr>
<tr>
<td>TT</td>
<td>43.5</td>
<td>25.7</td>
<td></td>
</tr>
<tr>
<td>MTHFR 1298A&gt;C</td>
<td>(%)</td>
<td>(%)</td>
<td></td>
</tr>
<tr>
<td>AA</td>
<td>77.3</td>
<td>81.1</td>
<td>0.69†</td>
</tr>
<tr>
<td>AC</td>
<td>22.7</td>
<td>18.9</td>
<td></td>
</tr>
</tbody>
</table>

* Based on Mann-Whitney tests  † Based on χ² tests  § Hardy-Weinberg equilibrium test among controls for MTHFR 677C>T and MTHFR 1298A>C >0.05  # 1 case had missing values for MTHFR 1298A>C  P<sub>10-90</sub>: Percentile 10 and 90

### Table III

**MTHFR 677C>T AND 1298A>C POLYMORPHISMS AND RISK OF SPONTANEOUS ABORTIONS IN A MEXICAN PERINATAL COHORT.**

**Morelos, Mexico, 2001-2004**

<table>
<thead>
<tr>
<th>Polymorphisms</th>
<th>All</th>
<th>Early Abortions (≤ 10 weeks)</th>
<th>Late Abortions (&gt;10 weeks)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>OR&lt;sup&gt;†&lt;/sup&gt; 95% CI</td>
<td>Cases</td>
<td>Controls</td>
</tr>
<tr>
<td>MTHFR 677C&gt;T&lt;sup&gt;§&lt;/sup&gt;</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CC+CT</td>
<td>1.0  --</td>
<td>5  55</td>
<td>1.0  --</td>
</tr>
<tr>
<td>TT</td>
<td>5.0 1.2  20.9</td>
<td>4  19</td>
<td>5.2 0.75  35.6</td>
</tr>
<tr>
<td>MTHFR 1298A&gt;C&lt;sup&gt;‡,§&lt;/sup&gt;</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>AA</td>
<td>1.0  --</td>
<td>8  60</td>
<td>--  --</td>
</tr>
<tr>
<td>AC</td>
<td>5.5 1.1  26.6</td>
<td>1  14</td>
<td>--  --</td>
</tr>
</tbody>
</table>

* Risk adjusted for maternal occupation, MTHFR 298 A>C, folate intake and caloric intake and husband's smoking status  † Risk adjusted for maternal occupation, MTHFR 677C>T, folate intake and caloric intake and husband's smoking status  § 1 case had missing values for the MTHFR 1298A>C
ing a longitudinal assessment of beta human chorionic gonadotrophin (ß-hCG),\(^4\) that was the case of this study and might have reduced the power to detect significant associations. Balancing these limitations is the strength of our study design, a nested case-control study, where dietary information and sera were obtained at recruitment and allow showing the temporality of the studied variables.

Maternal paid employment was associated with risk of SA in our study population. However, interpretation of this result is limited since we did not collect detailed work-related information such as the number of hours worked, type of exposures, etc. Our findings regarding paternal tobacco use are consistent with the study conducted by Venners et al.,\(^3\) which reported an almost twofold increase of SA among women whose husbands smoked more than 20 cigarettes per day, however the limited sample size precluded the ability to find statically significant results that should be replicated in future larger cohorts.

The mechanism by which the MTHFR polymorphisms increase the risk of SA is not known. Previously, authors have hypothesized that the increased risk may be due to an increase in homocysteine levels,\(^10,16\) which is consistent with the higher levels of homocysteine that we detected in our cases compared to the controls. Hyperhomocysteinemia reduces the methyl availability;\(^47\) methylation is an important epigenetic characteristic that plays an important role in DNA repair and the stability of the genome. Both MTHFR677 C>T and the MTHFR1298 A>C polymorphisms have been associated with decreased levels of DNA methylation.\(^17\) Experimental evidence also suggests that aberrant DNA methylation may contribute to the under-development observed in fetal cow clones.\(^48\)

Prevention of SA should consider metabolic dietary differences among the genetically susceptible population. Future larger cohort studies are needed to fully evaluate the interaction between the genetic characteristics and the periconception intakes of folate and other related B vitamins in high risk populations.

**Acknowledgements**

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**References**


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