

# The Association of Moderate/Severe Periodontitis (MSP), Pregnancy, and Two Single Nucleotide Polymorphisms (SNPs) in the Vitamin D Receptor (VDR) Gene: NHANES III, 1988-1994



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# Background

- Pregnancy can exacerbate an existing periodontal condition
- Periodontal disease has been associated with a number of adverse reproductive outcomes
- Activation of inflammatory pathway implicated in adverse reproductive outcomes since oral microorganisms and cytokines have been identified in amniotic fluid

## Purpose

To identify candidate genes for adverse reproductive outcomes using periodontal conditions as a "surrogate" outcome

## Objectives

- What genes are associated with periodontal conditions in a population of reproductive aged women?
- Can pregnancy modify this association?

### Methods

# Data Sources: NHANES III Collaborative Genomics Project

- 1. NHANES III (1988-1994)
  - National population-based weighted survey data (NCHS)
  - Over-sampling of youth, elderly, African-Americans, and Mexican-Americans
- 2. NHANES III DNA Data Bank (1991-1994)
  - Approximately 7,000 DNA samples >12 years of age
- 3. 2002 CDC Cross-CIO Collaborative Working Group

# Case/Control Study of Moderate/Severe Periodontitis (MSP)

- Candidate gene association study
- Study population of women 17-54 years of age (n = 2,103)
- Case definition of periodontitis based on CDC definition of moderate/severe periodontitis (MSP)

#### **Analysis**

- We used the log-additive model to identify genes associated with periodontitis among 13 genes/25 SNPs in the inflammatory metabolic pathways
- Multiple logistic regression was performed on significant associations using potential confounders as covariates
- Gene-environment interactions were included

### Results

#### Table 1

Study population characteristics among Non-Hispanic white (NHW), Non-Hispanic black (NHB), and Mexican-Americans (MA) (n = 2,103) Sample sizes are unweighted

Variable		% (SE)		p-value
	<u>NHW</u>	<u>NHB</u>	MA	
	n=679	n=763	n=661	
Mean Age (y)	34.7 (0.7)	33.0 (0.4)	32. 0 (0.4)	<0.001
Education (y)	, ,	, ,	,	
<12	12.4 (1.4)	22.7 (2.2)	54.2 (4.0)	
12	35.1 (2.2)	42.4 (2.6)	30.3 (3.0)	
>12	52.5 (2.7)	34.9 (2.9)	15.6 (2.3)	0.10
Tobacco Use	, ,	,	,	
Never smoker	53.9 (2.6)	66.0 (2.5)	75.2 (1.9)	0.02
Former smoker	18.9 (1.7)	7.6 (1.0)	13.0 (1.3)	
Current smoker	27.2 (2.2)	26.4 (2.0)	11.8 (1.4)	
Dental visit in the last 12	months	,	,	
Yes	60.0 (2.8)	46.9 (2.3)	36.1 (2.4)	0.01
No	40.0 (2.8)	53.1 (2.3)	63.9 (2.4)	
Poverty Income Ratio	, ,	,	,	
<100	8.3 (1.8)	36.6 (3.5)	37.1 (3.2)	
100-199		26.8 (2.2)	33.6 (2.8)	
>200	73.5 (3.0)	36.6 (3.2)	29.3 (1.5)	0.06
Diabetes	,	,	,	
Yes	2.7 (0.8)	4.4 (0.9)	6.6 (1.2)	
No	97.3 (0.8)	95.6 (0.9)	93.4 (1.2)	0.64
Reproductive History	,	, ,	,	
Never been pregnant	25.8 (2.3)	16.2 (1.6)	15.8 (2.6)	<0.001
Ever been pregnant	74.2 (2.3)	83.8 (1.6)	84.2 (2.6)	

#### Table 2

Prevalence of MSP and characteristics of the population with MSP by race/ethnicity (n= 94)

Variable		% (SE)		p-value
	<u>NHW</u>	<u>NHB</u>	MA	
Prevalence	2.5 (0.7)	6.1 (1.1)	3.5 (0.6)	0.05
Mean Age (y)	44.5 (1.3)	37.6 (1.3)	44.7 (1.6)	<0.001
Education (y)				
<12	22.5 (10.0)	38.2 (9.6)	78.2 (9.4)	0.14
12	35.5 (11.3)	32.2 (9.6)	18.4 (8.0)	
>12	41.9 (14.1)	29.1 (8.9)	3.4 (2.6)	
Tobacco Use	,	,	,	
Never smoker	14.8 (9.4)	53.7 (9.8)	64.3 (9.7)	0.008
Former smoker	26.2 (14.3)	3.9 (2.6)	10.5 (8.1)	
Current smoker	59.1 (16.5)	42.4 (Î0.1)	25.2 (9.1)	
Dental visit in the last 12	months	,	,	
Yes	25 (12.5)	47.5 (8.4)	26.8 (10.3)	0.02
No	75 (12.5)	52.5 (8.4)	73.2 (10.3)	
Poverty Income Ratio	,	,	` ,	
<100	27.6 (12.5)	27.7 (6.7)	53.5 (9.8)	
100-199	` '	36.9 (7.4)		
>200	48.0 (14.8)	36.5 (7.7)	17.6 (8.7)	0.2
Diabetes	,	,	, ,	
Yes	0 (0)	8.5 (4.6)	19.2 (8.0)	0.7
No	100 (0)	,	` '	
Reproductive History	( )		,	
Never been pregnant	0 (0)	12.1 (4.6)	3.9 (3.0)	0.001
Ever been pregnant	100 (0)	87.9 (4.6)	96.1 (3.0)	

# **Table 3**Genes/SNPs in the inflammatory/metabolic pathways associated with MSP (unadjusted)

`	,				
Gene	p-Value	Gene	p-Value	Gene	p-Value
CCR2	0.99	IL4R-05	0.58	MGC4093-03	0.03
CRP-B	0.12	-06	0.43	MGC4093-01	0.38
CRP-C	0.45	IL10-014	0.20	TNF-01	0.09
FCGR2A	0.57	-02	0.21	-04	0.32
IL1B-09	0.53	-03	0.30	-02	0.05
IL4 -01	0.90	MBL2 -11	0.02	MTHFR-01	0.57
-17	0.72	-12	0.74	-02	0.82
-02	0.54	TGFB1-01	0.16	VDR-01	0.009
				-07	0.003

#### Table 4

Allele frequencies of Vitamin D Receptor (VDR) SNP rs731236; C<T and rs2239185; C<T by case/control status (cases = with MSP, controls = no MSP) among NHW, NHB, and MA.

VDR SNP rs731236						
	1	<u>WHW</u>	1	NHB	1	MA
	Cases	Controls	Cases	Controls	Cases	Controls
C	61%	39%	35%	28%	34%	24%
Т	39%	61%	65%	72%	67%	76%

VDR SNP rs2239185						
	1	VHW		NHB	1	MA
	Cases	Controls	Cases	Controls	Cases	Controls
C	19%	349%	41.5%	42%	56%	58%
Т	81%	51%	58.5%	58%	44%	42%

#### Table 5

VDR SNP associations with MSP stratified by reproductive history and race/ethnicity (unadjusted)

VDR SNPs	rs731236	rs2239185
	unadjusted OR	unadjusted OR
	(95% CI)	(95% CI)
Reproductive History	<del>(2222-24</del>	<del>1</del>
Never been pregnant	0.5 (0.2-1.1)	0.5 (0.3-0.8)
Ever been pregnant	1.9 (1.2-2.9)	0.3 (0.1-0.8)
Race/ethnicity		
NHW	2.3 (1.2-4.5)	0.3 (0.1-0.8)
NHB	1.4 (0.9-2.2)	1.0 (0.6-1.5)
MA	1.6 (0.9-3.1)	0.9 (0.6-1.6)
IVI <i>/</i> =\	1.0 (0.3-3.1)	U.3 (U.U-1.U)

#### Table 6

VDR SNP associations with MSP (adjusted)

VDR SNPs	rs731236	rs2239185
Mean Age (y)	<u>adjusted OR</u> (95% CI) 1.1 (1.1 – 1.2)	<u>adjusted OF</u> (95% CI) 1.1 (1.1 – 1.2)
Race/ethnicity NHW	REF	REF
NHB	4.3 (0.7 – 28.3)	0.2 (0.03 - 1.3)
MA	2.9 (0.3 – 27.0)	0.06 (0.01 – 0.7
Tobacco Use		
Never smoker	REF	REF
Former smoker	1.8 (0.6 - 6.0)	2.2 (0.6 - 8.4)
Current smoker	4.1 (1.2 – 14.2)	4.6 (1.2 – 17.0)
Reproductive History		
Never been pregnant	REF	REF
Ever been pregnant	0.1 (0.01 – 1.3)	1.6 (0.1 – 22.5)
VDR	0.5(0.2-1.4)	0.2 (0.1 – 0.6)
VDR* Reproductive History		
Never been pregnant	REF	REF
Ever been pregnant	5.7 (1.6 – 20.5)	1.2(0.3-5.4)
VDR* Race/ethnicity		
NHW	REF	REF
NHB	0.8 (0.3 - 1.7)	3.4 (1.1 – 10.4)
MA	0.6(0.2-2.0)	4.8 (1.3 – 17.1)

\* Adjusted for education, number of dental visits, poverty income ratio, & diabetes status

# **Summary and Limitations**

#### Summary

- Identified two SNPs in the VDR gene to be associated with MSP
- Women who were ever pregnant and who had MSP were more likely to be carriers of rs731236 compared to non-pregnant women
- Non-Hispanic black women and Mexican American women were more likely to carry VDR SNP rs2239185 compared to non-Hispanic white women

# Study limitations

- Small number of MSP cases in each race/ethnicity category
- All non-Hispanic white women who had MSP also had a past pregnancy