Junil FOR RESERANCE	Original Research Paper	Orthopedics		
And the matter of the matter o	PLASMA VITAMIN D LEVEL AND VITAMIN D RECEPTOR GENETIC POLYMORPHISM OF SPINAL TUBERCULOSIS SUSCEPTIBILITY IN WEST JAVA POPULATION, INDONESIA: DOES IT CORRELATED?			
Ahmad Ramdan	Department of Orthopaedic & Traumato Universitas Padjadjaran, Bandung	logy, Faculty of Medicine,		
Ida Parwati	Department of Clinical Pathology, Faculty of Medicine, Universitas Padjadjaran, Bandung			
Budi Setiabudiawan	Department of Paediatrics, Faculty o Padjadjaran, Bandung	of Medicine, Universitas		
Bambang Tiksnadi	Department of Orthopaedic & Traumato Universitas Padjadjaran, Bandung	logy, Faculty of Medicine,		
ABSTRACT To explores the potential role of plasma vitamin D level and VDR gene polymorphisms in susceptibility to tuberculo				

ABSTRACT is optioned to be obtained of plasma vitamin D level and VDR gene polymorphisms in susceptibility to tuberculo sis (TB) in the West Java population, Indonesia. A case-control study was performed on 53 Spinal TB patients, 53 Pulmonary TB patients and 53 healthy controls. Plasma Vitamin D levels were analyzed by ELISA, genetic polymorphisms of VDR gene (Taql, Fokl, Bsml, Apal) were analyzed using PCR-RFLP and sequencing, There was a significant difference (p<0.01) of plasma vitamin D level among all groups. Genotype analysis of VDR gene polymorphismFok1, Bsml, Apal and Taql has shown no significant difference in all group (p>0.01). The combination of vitamin D deficiency and VDR gene polymorphism Fok1 in FF genotype were contributing factors to the susceptibility of spinal tuberculosis in the West Java population, Indonesia.

KEYWORDS : Spinal tuberculosis, Vitamin D, VDR gene polymorphism

# INTRODUCTION

*M. tuberculosis* infection of the musculoskeletal system, especially the spine, is caused by differences in the immune response to tuberculosis. Many studies suggest that the occurrence of TB is not only determined by bacterial factors alone, but also by the body immune system mechanisms that suppress *M. tuberculosis* virulence. <sup>1,2</sup> Lack of vitamin D (1,25-*dihydroxycholecalciferol*) is associated with the risk of TB infection. People with vitamin D deficiency have a greater susceptibility to infections of tuberculosis.<sup>3,5</sup>

Previous studies showed that abnormalities of vitamin D receptor structure, in this case polymorphism, has a role in immune system activity against TB infection. Some types of vitamin D receptor polymorphisms gene such as *Bsm1*, *Apa1*, *Taq1* and *Fok1* are thought to correlate with the incidence of TB.<sup>612</sup>

There have been no studies done in Indonesia about the correlation of plasma vitamin D level and the role of vitamin D receptor gene *Bsm1*, *Apa1*, *Taq1* and *Fok1*. This encourage researcher to analyze the role of vitamin D levels in plasma and vitamin D receptor polymorphism gene *Bsm1*, *Apa1*, *Taq1* and *Fok1* against immunopathology that affect the vulnerability and resilience of spinalTB infection.

# METHOD

This is case control study. The research subjects are spinal tuberculosis that met the inclusion criteria, namely: spinal tuberculosis patients that had been operated, pulmonary tuberculosis patients and healthy patients with age  $\geq$  15 years old. Cases group consist of 53 spinal TB patients, 53 pulmonary tuberculosis patients and control groups consist of 53 healthy patients. Total plasma vitamin D measured for all sample by ELISA. Polymorphism of vitamin D receptor (VDR) done by sequencing method and PCR-RFLP technique. All statistical analysis was processed using SPSS version 16. The Chi-square test was used to compare the frequency of variable, p< 0,05 was considered statistically significant.

# RESULTS

Based on Table 1, there were significant differences of plasma vitamin D level in all three groups (p <0.01). In further comparison test, it was showed that plasma vitamin D concentrations and tuberculosis was closely related (mean differences of normal and spinal TB patients = 4, 35 and mean differences of normal and pulmonary TB patients = 4, 19; p = 0,000). PCR products restricted with Fok1 reveals genotypes (after treatment with enzyme) denoted FF (265 bp), Ff (265,196 and 69 bp), or ff (196 and 69 bp); Bsm1 genotypes denoted B8 (825 bp), Bb (825, 650 and 175 bp), bb (650 and 175 bp); Apa1 genotypes denoted AA (740 bp), Aa (740, 530 and 210 bp), aa (530 and 210 bp) and Taq1 genotypes denoted TT (495 and 245 bp), Tt (495, 290, 245 And 205 bp), tt (290, 245 and 205 bp).

Vitamin D	Normal (n=53)		nal TB =53)	Pulmonar y (n=53)	p value
Plasma Vitamin D Level					
Mean (SD)	11,90 (6,83)	7,55	6,32)	7,71 (8,06)	0,000**a)
Median	10,82	5,53		4	
(Range)	(3 -33,88)	(3 -	28,21)	(3 - 41,77)	
Comparison		p-value			
Normal vs Spinal TB		0,000b			
Normal vs Pulmonary TB		0,000b			
Spinal TB vs Pulmonary TB		0,445			

TABLE 1- Analysis and Further Comparison Test Plasma Vitamin D Level

p value was measured from a) Kruskal Wallis(data not normally distributed), b), P value was measured from Mann Whitney test, difference consideredc significantif p < 0,05 and covery significantif p < 0,01.

Allele and genotype frequencies of VDR gene polymorphisms in the spinal TB, pulmonary TB and normal group were analyzed by SPSS 16 program as shown in Table 2. Table 3 showed, all genotype frequencies of VDR gene were not a significant association with plasma Vitamin D level concentration except Fokl-FF genotype (p=0.030).

TABLE 2- Proportional Analysis of Genotype dan AlleleVDR gene polymorphismin Spinal TB group, Pulmonary TB group and case control group

Spinal TB         Pulmonary TB           Spinal TB         TB           (n=53)         (n=53)           Bsm1	<ul> <li>control</li> <li>(n=53)</li> <li>3 (5,66)</li> <li>5 (9,43)</li> <li>45 (84,91)</li> <li>8 (15,09)</li> <li>45 (84,91)</li> </ul>	Value 0,562 0,810
(n=53)         (n=53)           Bsm1         (m=53)           BB         0 (0)         2 (3,77)           Bb         6 (11,32)         6 (11,32)           bb         47 (88,68)         45 (84,91)           Alleles         (m=53)         (m=53)           B         6 (11,32)         6 (11,32)           bb         47 (88,68)         45 (84,91)           Alleles         (m=53)         (m=53)           A         6 (11,32)         8 (15,09)           b         47 (88,68)         45 (84,91)           Apa1         (m=53)         (m=53)           AA         3 (5,66)         4 (7,55)           Aa         24 (45,28)         23 (43,4)           aa         26 (49,06)         26 (49,06)           Alleles         (m=53)         (m=53)           A         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           Taq1         (m=57)         49 (92,45)	(n=53) 3 (5,66) 5 (9,43) 45 (84,91) 8 (15,09)	
Bsm1         Image: state of the state	3 (5,66) 5 (9,43) 45 (84,91) 8 (15,09)	
BB         0 (0)         2 (3,77)           Bb         6 (11,32)         6 (11,32)           bb         47 (88,68)         45 (84,91)           Alleles	5 (9,43) 45 (84,91) 8 (15,09)	
Bb         6 (11,32)         6 (11,32)           bb         47 (88,68)         45 (84,91)           Alleles             B         6 (11,32)         8 (15,09)           b         47 (88,68)         45 (84,91)           Apa1             AA         3 (5,66)         4 (7,55)           Aa         24 (45,28)         23 (43,4)           aa         26 (49,06)         26 (49,06)           Alleles             A         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           Taq1             TT         48 (90,57)         49 (92,45)	5 (9,43) 45 (84,91) 8 (15,09)	
bb         47 (88,68)         45 (84,91)           Alleles	45 (84,91) 8 (15,09)	0.810
Alleles         (4,7,7)           B         6 (11,32)         8 (15,09)           b         47 (88,68)         45 (84,91)           Apa1	8 (15,09)	0.810
B         6 (11,32)         8 (15,09)           b         47 (88,68)         45 (84,91)           Apa1		0.810
b         47 (88,68)         45 (84,91)           Apa1         -         -           AA         3 (5,66)         4 (7,55)           Aa         24 (45,28)         23 (43,4)           aa         26 (49,06)         26 (49,06)           Alleles         -         -           A         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           Taq1         -         -           TT         48 (90,57)         49 (92,45)		0.810
Apa1         (4,7,55)           AA         3 (5,66)         4 (7,55)           Aa         24 (45,28)         23 (43,4)           aa         26 (49,06)         26 (49,06)           Alleles	45 (84,91)	0,010
AA         3 (5,66)         4 (7,55)           Aa         24 (45,28)         23 (43,4)           aa         26 (49,06)         26 (49,06)           Alleles         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           A         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           Taq1		1
Aa         24 (45,28)         23 (43,4)           aa         26 (49,06)         26 (49,06)           Alleles         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           a         26 (49,06)         26 (49,06)           Taq1		
aa         26 (49,06)         26 (49,06)           Alleles         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           Taq1	9 (16,98)	0,305
Alleles         27 (50,94)         27 (50,94)           A         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           Taq1         71         48 (90,57)           TT         48 (90,57)         49 (92,45)	18 (33,96)	1
A         27 (50,94)         27 (50,94)           a         26 (49,06)         26 (49,06)           Taq1	26 (49,06)	1
a         26 (49,06)         26 (49,06)           Taq1		
Taq1         48 (90,57)         49 (92,45)	27 (50,94)	1,000
TT 48 (90,57) 49 (92,45)	26 (49,06)	1
	47 (88,68)	0,702
Tt 5 (9,43) 4 (7,55)	5 (9,43)	1
tt 0 (0) 0 (0)	1 (1,89)	1
Alleles		
T 53 (100) 53 (100)	52 (98,11)	0,366
t 0 (0) 0 (0)	1 (1,89)	1
Fok1		
FF 14 (26,42) 22 (41,51)	20 (37,74)	0,351
Ff 27 (50,94) 23 (43,4)	27 (50,94)	]
ff 12 (22,64) 8 (15,09)	6 (11,32)	1
Alleles		
F 41 (77,36) 45 (84,91)	47 (88,68)	0,276
f 12 (22,64) 8 (15,09)	6 (11,32)	]

P value was measured from Chi Square test, difference considered \*significant if p < 0,05 and \*\*very significant if p < 0,01

### TABLE 3- Correlation Analysis Between Genotype VDR Gene Polymorphism with Mean Plasma Vitamin D Level in All Group

	Genotype	Plasma Vitamin D Level	
		Mean (SD)	P value
Bsm1	BB	14,24 (14,13)	0,197
	Bb	8,17 (5,13)	
	bb	8,93 (7,15)	
Apa1	AA	12,44 (11,21)	0,079
	Aa	7,96 (5,64)	
	aa	9,21 (7,41)	
Taq1	TT	9,2 (7,59)	0,578
	Tt	7,16 (3,84)	
	tt	12,24 (-)	
Fok1	FF	9,73 (7,41)	0,030*
	Ff	9,70 (7,99)	
	ff	5,50 (3,13)	

### DISCUSSION

In human, mycobacterial pathogenicity varies from one mycobacterial species to another. There is much variability among individuals in the response to mycobacterial infections, but it is not known why certain people develop disease when challenged with mycobacteria and others remain healthy.<sup>13</sup> Plasma vitamin D levels and the polymorphisms in VDR gene have been studied in relation to tuberculosis in many populations. Epidemiological evidence suggests that there is a link between vitamin D deficiency and susceptibility to leprosy and TB. Recently genetic variation has been shown to be associated with single nucleotide polymorphisms (SNP) in the vitamin D receptor gene in many populations of leprosy and TB.13-1

Vitamin D works in the immune system by lowering the levels of

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inflammatory proteins and increase the antimicrobial protein that can destroy bacteria such as tuberculosis. Vitamin D metabolism leads to activation of macrophages and restricts the intracellular growth of mycobacteria. This effect of vitamin D may be influenced by polymorphisms at three sites (Taq1, Bsm1 and Fok1) in the vitamin D receptor gene.<sup>13</sup> Previous studies show that people with low levels of vitamin D are more susceptible to TB than people with high vitamin D levels. Therefore it is clear that on our study (table 1) there are significant differences between the vitamin D levels in the normal group and the case group (spinal TB and pulmonary TB).Vitamin D deficiency causes secondary hyperparathyroidism and bone matrix and mineral loss and increased risk of osteoporosis and bone damage. In infections of the spine, bone destruction occurs which will aggravate the state of vitamin D deficiency.<sup>14</sup> This causes plasma vitamin D level in a group of spinal TB group (7,55) was inferior to pulmonary TB group (7,71) for the event of TB infection in the bone of vitamin D (table 1).

In our study, there was no statistically meaningful relationship between the allele and genotype frequencies of Fok1, Bsm1, Apa1 and Taq1 polymorphisms in the VDR gene and susceptibility to tuberculosis (table 2), but when they were analyzed with respect to plasma vitamin D concentrations, a significant association was seen (table 3). There was significant relationship in plasma vitamin D level with *Fok1*-FF genotype (p=0,030).

The findings in our study indicated that vitamin D deficiency ( $\leq 10$ ng/ml) (especially in Fok1-FF individuals) can carry a higher risk of tuberculosis; therefore moderate supplementation may be useful. Although several investigators reported and suggested that the VDR polymorphism may be of immunoregulatory importance for many disease processes, it is not clear that the polymorphism determine susceptibility to the development of clinical disease or susceptibility to infection. Further studies will be required to investigate how VDR polymorphism may influence susceptibility to infectious disease or development of clinical disease.<sup>1</sup>

# CONCLUSION

Deficiency of vitamin D plasma levels and receptor gene polymorphisms Fokl vitamin D (RVD) is a risk factor for spinal tuberculosis. The combination of plasma levels of vitamin D deficiency and homozygous genotypes of polymorphism Fokl FF RVD has the great chance of susceptibility of spinal tuberculosis in The West Java population, Indonesia.

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