Vitamin D receptor genetic polymorphisms and the risk of multiple sclerosis: A systematic review and meta-analysis

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There are conflicting results regarding the exact effect of the vitamin D receptor (VDR) gene polymorphisms on the susceptibility to multiple sclerosis (MS). Therefore, we aimed to investigate the impact of four major studied VDR gene polymorphisms consisting of Apal, Bsml, Fokl, and Tagl on the risk of MS in the Iranian population. A literature search was performed in various databases to find case-control studies evaluating the association between VDR gene polymorphisms and MS risk in Iran. Data were extracted and odds ratios (OR) with 95% confidence intervals (CI) were calculated. Subgroup analyze was performed to detect potential sources of heterogeneity. A total of 1206 cases and 1402 controls in nine case-control studies were included. Apal was the only variant which showed statistically significant relation in allelic (OR = 0.54 (95% CI: 0.37?0.79); P = 0.00), homozygote (OR = 3.48 (95% CI: 1.7?6.9); P = 0.00), dominant (OR = 0.56 (95% CI: 0.3?0.79); P = 0.01), and recessive (OR = 0.35 (95% CI: 0.18?0.66); P = 0.00) models. The Tagl polymorphism showed a significant negative association with MS only in the homozygote model (OR = 0.28 (95%) CI: 0.08?0.9); P = 0.04). The Bsml polymorphism also showed significant relation in allelic (OR = 0.69 (95% CI: 0.51?0.94); P = 0.01), homozygote (OR = 0.46 (95% CI: 0.25?0.86); P = 0.01), and recessive OR = 0.56 (95% CI: 0.39?0.8); P = 0.00) models after performing sensitivity analysis. Fokl polymorphism showed no significant association with MS risk. Apal and Tagl TT genotype were found contributing to MS susceptibility and Bsml and Fokl showed no relation with MS susceptibility

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Meta-analysis
Multiple sclerosis
Vitamin D receptor
vitamin D receptor
allele
data base
data extraction
evaluation study
genetic heterogeneity
genetic polymorphism
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homozygote
human
Iranian people
meta analysis
multiple sclerosis
odds ratio
Review
risk assessment
sensitivity analysis
systematic review