



POSTER PRESENTATION

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# PreS-FINAL-2053: Vitamin D receptor polymorphisms in a cohort of Italian patients with juvenile idiopathic arthritis

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## Introduction

A role for vitamin D has been hypothesized in generating disease activity for patients with juvenile idiopathic arthritis (JIA): specific polymorphisms of vitamin D receptor (*VDR*) gene have recently been associated with different biologic response to vitamin D itself.

## Objectives

To evaluate *VDR* polymorphisms in patients with JIA in comparison with unrelated healthy controls.

## Methods

We recruited 63 Italian children, adolescents and young adults with JIA (mean age 16.21 + 7.11 SD yrs, 51 female and 12 males, female/male ratio 4.25 from 1 Unit of Paediatric Rheumatology and 1 Unit of Rheumatology, Transition Clinic. After informed consent, during routine laboratory tests, their genomic DNA was extracted from peripheral blood leukocytes, to analyze *VDR* polymorphisms by PCR-based sequencing (*CDX2* in the promoter region) and PCR-based enzymatic digestions (*FokI* in exon 2, *BsmI* and *ApaI* in intron 8, and *TaqI* in exon 9). An Italian population of 2221 unrelated individuals without JIA was used as healthy controls.

## Results

The distribution of *FokI*, *BsmI*, *ApaI*, and *TaqI* polymorphisms did not show significant differences between children with JIA and controls. Regarding the *CDX2* polymorphism, we observed a statistical difference in the distribution of GG and GA genotypes, with the GG genotype more frequent in JIA subjects (Yates-corrected chi-square

6.97; Odds ratio = 2.08;  $p = 0.008$ ) and the GA genotype in healthy controls (Yates-corrected chi-square 4.04; Odds ratio = 0.55;  $p = 0.044$ ). Data about AA genotype were not significant due to their very low number (three) within the JIA population. G allele resulted to be more frequent in JIA subjects (Yates-corrected chi-square 6.51; Odds ratio = 1.82;  $p = 0.011$ ).

## Conclusion

Pathogenetic mechanisms influencing the predisposition to JIA are poorly elucidated. Our analysis of *CDX2* polymorphisms located in the promoter region of the *VDR* gene has revealed that both GG genotype and G allele are more represented in patients with JIA. By our preliminary data, we can speculate that the G allele decreases *VDR* transcriptional activity with respect to the A allele, as well as the presence of GG genotype could explain a reduction of *VDR* activity, with subsequent decreased response to vitamin D and potential immunity deregulation leading to JIA.

## Disclosure of interest

None declared.

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