

The Effect of SNPs and Interferon Therapy on *PSMC6* Gene Expression in Multiple Sclerosis Patients in Latvia



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Background

Multiple sclerosis (MS) is a neurodegenerative and autoimmune inflammatory disease that leads to various neurological disabilities [1]. Ubiquitin-proteasome system (UPS) is a key player in degradation of damaged and unnecessary proteins. Dysfunction of ubiquitin-proteasome system (UPS) causes accumulation of toxic protein aggregates in MS patients [2]. For treatment of MS, the first class of disease modifying therapies is interferon β (IFN β). It interacts with the immune system: B cells, T cells, antigen-presenting cells, thus providing clinical efficiency for MS patients [3].

Aim

The aim of the current study was to evaluate SNPs of *PSMC6* gene as possible biomarkers for multiple sclerosis in Latvian population.

Methods

Altogether 54 MS patients treated with IFN β therapy and 44 MS patients without the treatment were enrolled in the study. RNA was isolated from blood and *PSMC6* gene expression was analysed with qPCR. The *PSMA6* rs2295826 and rs2295827 were genotyped on MS subtype-, sex- and treatment efficiency association in 280 cases /305 controls study.

Kolmogorov-Smirnov (if $N > 50$) or Shapiro-Wilk (if $N < 50$) tests were used to test the normality. Comparisons between groups were done with Mann Whitney U test for two groups or Kruskal-Wallis test for more than two groups. Association with genotyping results was calculated using eta (η).

References

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2. David *et al.* Detection of protein aggregates in brain and cerebrospinal fluid derived from multiple sclerosis patients. *Front Neurol.* 2014;5:251.
3. Kasper *et al.* Immunomodulatory activity of interferon-beta. *Ann Clin Transl Neurol.* 2014;1(8):622-31.

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Results

In the group of MS patients treated with IFN β , patients with AA/CC genotype showed statistically significantly higher *PSMC6* gene expression compared to patients with AG/CT+GG/TT genotypes. When all of the groups were analysed together (according to genotypes and treatment), the results showed statistically significant differences among the groups (Fig. 1).

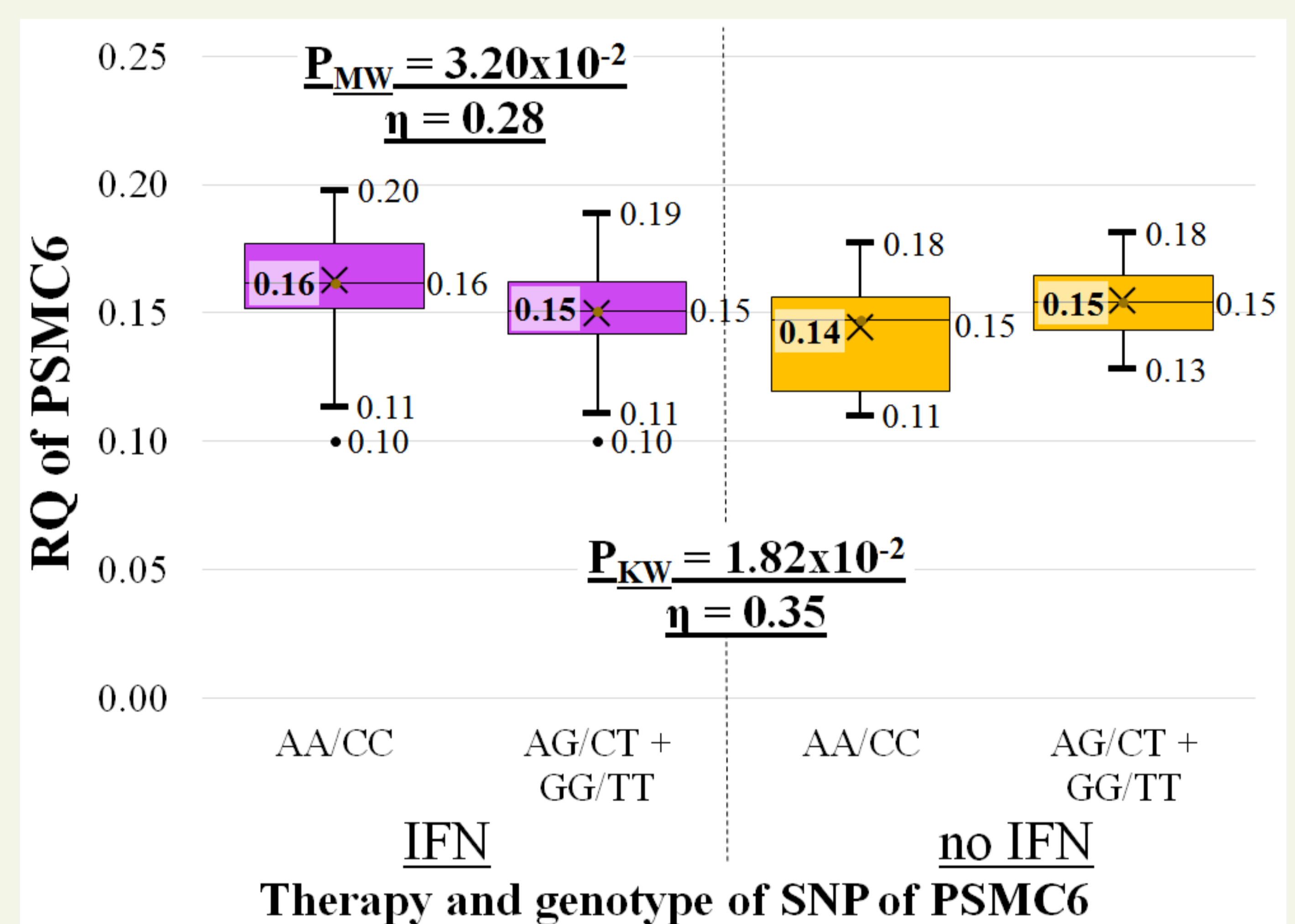


Figure 1. Expression levels of *PSMC6* gene in multiple sclerosis (MS) patients. IFN – group of MS patients with interferon β treatment; no IFN – group of MS patients without the treatment. AA/CC – common genotype; AG/CT+GG/TT – rare alleles and heterozygous genotypes.

Conclusions

The results indicate that both the treatment with IFN β and genotypes of patients affect the expression of *PSMC6* gene.