

THE ROLE OF hsp70-2, hsp70-hom HEAT SHOCK PROTEINS ON MULTIPLE SCLEROSIS RISK AND SEVERITY

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Introduction

Oxidative stress is involved in MS pathogenesis and progression by direct and indirect mechanisms of action [1], which also encompass the heat shock proteins 70 (hsp70s) family [2].

Intracellular hsp70s act as chaperones and anti-apoptotic proteins[3]. Extracellular hsp70s process and present antigens, promoting the activation of immune system [4]. Polymorphisms leading to either quantitative or qualitative change in hsp70 expression likely affect both the hsp70 cyto-protective and/or immune-modulatory effects.

Among the several proteins included in hsp70s family, the two major stress-inducible members (i.e. the Hsp70-1 and Hsp70-2) are encoded by HSPA1A and HSPA1B gene respectively, and the constitutively expressed non-inducible protein (i.e. Hsp70-hom) is encoded by HSPA1L gene. These three genes are located on chromosome 6 (6p21.3)[5], within the human leukocyte antigen (HLA) class III region.

While polymorphisms within HSPA1A exons are silent, we recently demonstrated that +1267 A/G HSPA1B (rs1061581) polymorphism is associated with an increased MS risk and that MS patients with GG or GA genotype display a significant reduction of hsp70-2 expression compared to patients with AA genotype[6]. Polymorphisms in the HSPA1L gene are mainly located in the region coding for the substrate-binding domain.

Aims

To investigate the association of MS with HSP70-hom polymorphism and haplotypes including another HSP70 gene known as HSP70-2 and to evaluate the relationship between Hsp70-hom protein expression level and MS severity, using peripheral blood mononuclear cells (PBMC) from MS patients and healthy donors.

Methods

- We included 195 MS Caucasian patients from the MS Centre of the National Neurological Institute "C. Mondino" (Pavia, Italy) and 439 Caucasian Healthy Controls.
- HSP70-hom polymorphism was studied with PCR-RFLP. Western blot analyses were performed to quantify the Hsp70-hom protein expression levels in PBMC.
- We performed additive and genotypic unconditional logistic regression analyses, sex and age adjusted, to assess the association between the HSP70 polymorphisms and MS risk.
- The quantitative expression of the protein was tested using linear regression models, sex and age adjusted.
- We also tested whether the overall HSP70-2 and HSP70-hom haplotype variation influences MS presence and severity (quantified by multiple sclerosis severity score, MSSS [7], performing omnibus and conditional LR-based tests (*haplotype-based association testing tool, plink 1.07*).

Results

- The minor allele C of rs2227956 (HSP70-hom) conferred a risk against the MS (OR=2.13, P<0.0001) and having the CC genotype compared to TT genotype increased the risk of the MS almost 7 times (OR=6.71, P<0.0001)
- MSSS score was increased of 1.21 (P=0.017) among CC carriers (Figure 2) w.r.t. TT carriers
- The frequency of HSP70-2 and HSP70-hom GC haplotype (risk alleles combination) in MS patients respect to controls was significantly increased (OR: 3.489, p-value <0.0001) Intriguingly, HSP70-hom showed independent effect on MS risk after adjusting for the effect of the overall haplotype configuration (conditional LR test p-value<0.0001)
- The Hsp70-hom protein expression in PBMC of 47 MS patients and 37 controls were not significantly different by HSP70-hom genotype
- We explored the possible relationship between HSP70-hom protein low expression and mild MS : HSP70-hom protein total amount in low-severity patients (MSSS score <3, 1069±1035, median: 676 arbitrary units) was significantly lower vs. high-severity patients (MSSS score ≥3, 1730±1536, median: 1134 arbitrary units), Wilcoxon rank-sum test, P=0.038 (Figure 4).

Discussion and Conclusions

- We observed an increased risk of MS in HSP70-hom rs2227956 C carriers and the reduced expression of hsp70-hom in MS patients with a mild form of the disease
- HSP70-hom plays a more relevant role in promoting a pro-inflammatory immune system activation and an effective T cell response against the myelin antigens compared with its role in protecting CNS cells from inflammatory injury
- However, the underlying mechanisms involved in this unfavourable outcome are not clear yet and further studies are required to clarify the exact roles of hsp70-hom and its possible applications as biomarker and/or as target therapy in MS

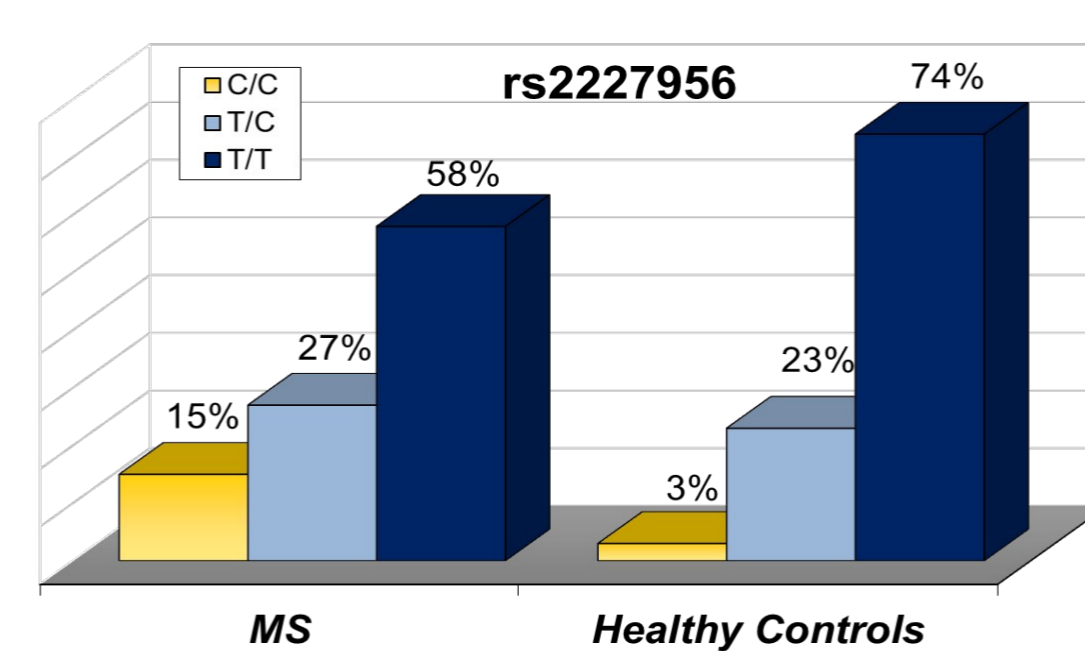


FIGURE 1. HSP70-hom polymorphism frequency distribution among MS cases and controls and MS estimates of risk among mutated genotypes

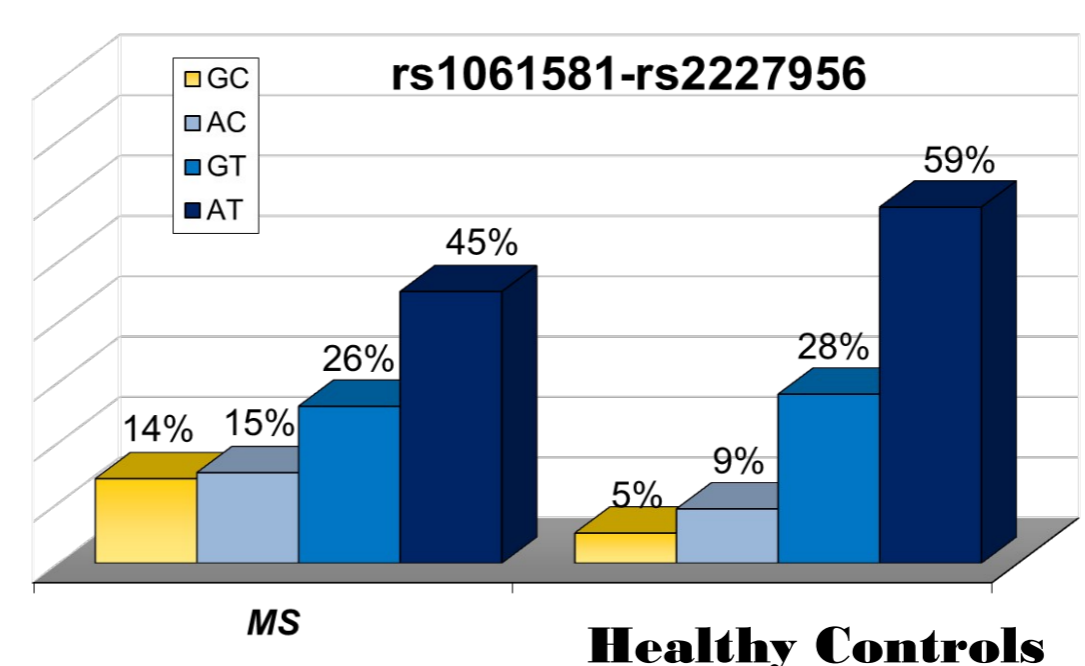
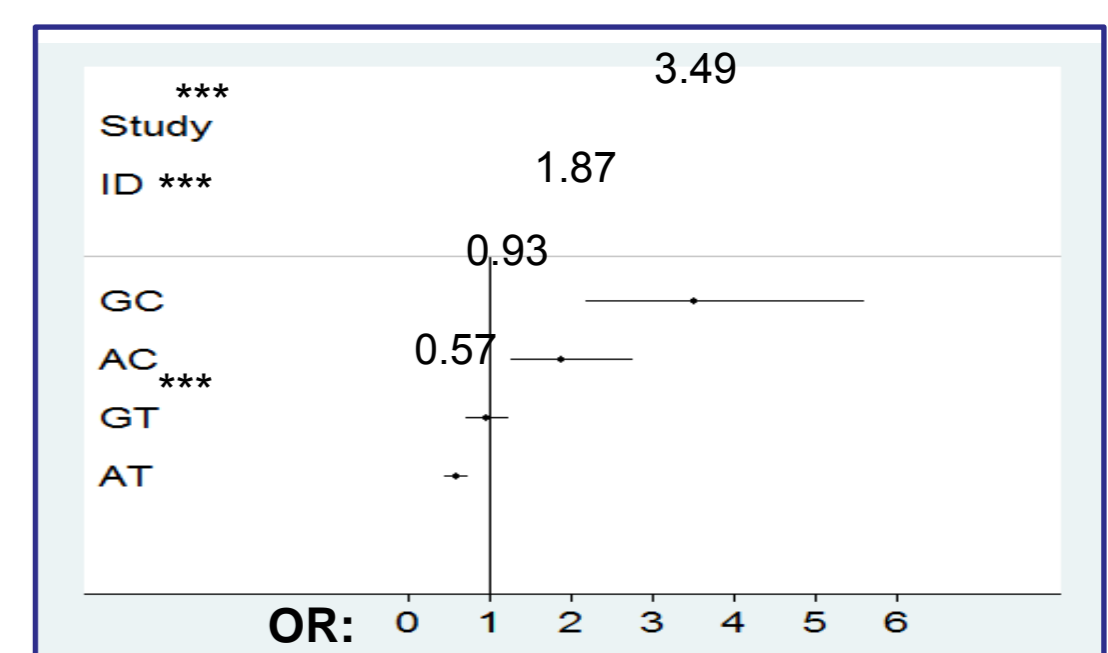
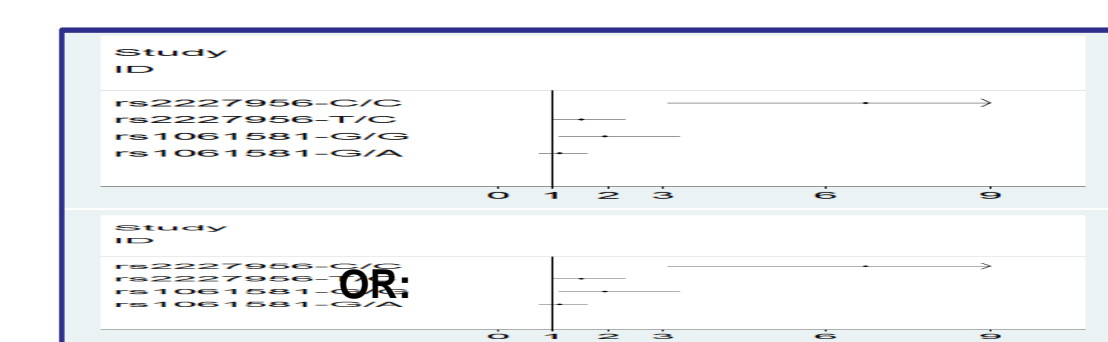


FIGURE 2. HSP70-2/HSP70-hom haplotypes frequency distribution among MS cases and controls and MS estimates of risk



HEALTHY CONTROLS

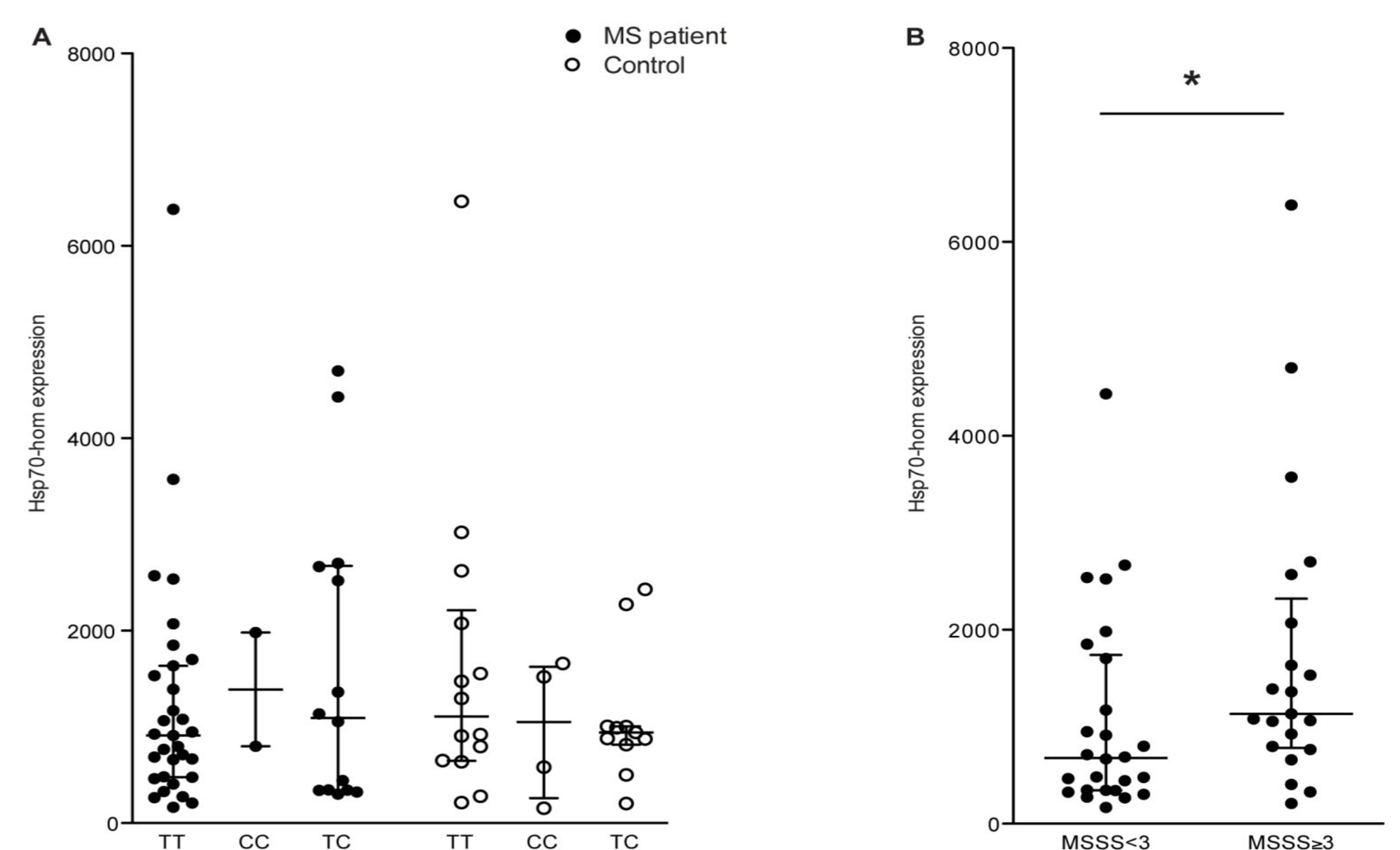


Figure 3

FIGURE 3. HSP70-hom expression by genotype and MS impact

References

- Lassmann H. Mechanisms of white matter damage in multiple sclerosis. *Glia*. 2014 62: 1816-1830.
- Mansilla MJ, Montalban X, Espejo C. Heat shock protein 70: roles in multiple sclerosis. *Mol Med*. 2012 18: 1018-1028.
- Mayer MP. Hsp70 chaperone dynamics and molecular mechanism. *Trends Biochem Sci*. 2013 38: 507-514.
- Li Z, Menoret A, Srivastava P. Roles of heat-shock proteins in antigen presentation and cross-presentation. *Curr Opin Immunol*. 2002 14: 45-51.
- Brocchieri L, Conway de Macario E, Macario AJ. hsp70 genes in the human genome: Conservation and differentiation patterns predict a wide array of overlapping and specialized functions. *BMC Evol Biol*. 2008 8: 19.
- Boiocchi C, Oserra C, Monti MC, et al. Are Hsp70 protein expression and genetic polymorphism implicated in multiple sclerosis inflammation? *J Neuroimmunol*. 2014 268: 84-88.
- Roxburgh RH, Seaman SR, Masterman T, et al. Multiple Sclerosis Severity Score: using disability and disease duration to rate disease severity. *Neurology*. 2005 64: 1144-1151