Letter

**CTLA4 polymorphism and primary Sjögren's syndrome**

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Similar to our study, the Gottenberg study analysed both CTLA4 CT60 and +49A/G single nucleotide polymorphisms (SNPs). They observed an association with the +49A/G SNP in one pSS cohort; however, this failed to replicate in a second cohort. We unreservedly agree with the authors’ conclusions regarding the importance of replication cohorts. However, we disagree that their results can be interpreted meaningfully against our study.

The CTLA4 CT60 and +49A/G SNPs are haplotype tags for three common, extended CTLA4 haplotypes [3,4], and the main finding of our study was an association between the +49A:CT60G haplotype and autoantibody positive pSS. As we noted in our paper, the individual SNPs occur on multiple haplotypes, and an individual SNP analysis, such as that reported by Gottenberg and colleagues, may result in negative studies or inconsistent findings between studies, particularly if the haplotype frequencies vary between study populations.

In addition to our finding of an association with pSS, the CTLA4 +49A:CT60G haplotype is also associated with systemic lupus erythematosus [5], which shares a number of clinical, serological and genetic features with pSS. A more recent study has identified several haplotype blocks across the extended CD28-CTLA4-ICOS region, with systemic lupus erythematosus associations observed in the distal 3’ flanking region of CTLA4 on a haplotype that includes variants in the promoter of ICOS [6]. Therefore, the balance of evidence supports a genetic association between this region and systemic autoimmune disease, although the precise nature, definition and boundaries of the haplotypes involved remain to be fully defined.

**Competing interests**
The authors declare that they have no competing interests.

**References**


pSS = primary Sjögren’s syndrome; SNP = single nucleotide polymorphism.