Impact of genetic predisposition of de novo uveitis with etanercept in ankylosing spondylitis

We read with interest the paper by Sieper et al reporting acute anterior uveitis (AAU) during etanercept or sulfasalazine therapies for axial spondyloarthritis (AxSpA). AAU is a well-known symptom in AxSpA without therapy and is itself a good indication for tumour necrosis factor (TNF) blockers. However during the last years, new onsets of AAU were reported with TNF blockers, as a paradoxical effect of this therapy, similarly to many cases reported. AAU with etanercept therapy, whereas rheumatic symptoms were under control, similarly to many cases reported. AAU resolved rapidly with local therapy without etanercept discontinuation in both patients. However, AAU relapsed and the decision was taken to switch etanercept to another TNF blocker: infliximab for the first case (who had been treated by adalimumab before) and adalimumab for the second case, with no AAU relapse to date. AAU was more frequently reported with etanercept, the soluble receptor p55, than with monoclonal antibody TNF blockers. This discrepancy between TNF blockers may be due to the specific structure of etanercept, which is a fusion protein combining two extracellular ligand-binding portions of the human soluble TNF receptor p75 with the end Fc fragment of human immunoglobulin G1.

Since de novo uveitis is a rare event with approximately 1 case for 100 patient-years exposed to TNF blockers, occurrence in two cousins from the same family, both treated for AxSpA by etanercept, suggested a genetic predisposition to this paradoxical side effect with TNF blockers. Since the microsatellite, D9S157, on chromosome 9p was identified by genome-wide scan to be associated with AAU in AxSpA, we analysed this region using several microsatellites in these two patients. Both patients shared an identical allele at three contiguous polymorphic markers, located at 9p21.3–p22.2: D9S157, D9S1684 and D9S162, but are different for D9S156 and D9S171. This locus covers approximately a 6 Mb interval and contains some 37 genes. This identity might be an additional indication for a genetic predisposition to uveitis at this locus, and along these lines, it is noteworthy that the cluster of interferon type 1 genes is present in this region between markers D9S162 and D9S171. Conversely, in our family, there is a probability of 1/16 that this haplotype identity occurred by chance. Further experiments are required to determine the specific gene from this area related to AAU.

According to our experience and the findings of this genetic analysis, we would suggest avoiding etanercept among TNF blockers when patients present a familial history of uveitis.

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