Genetic Variants in Epigenetics Factors and Rheumatoid Arthritis

Comprehensive pharmacogenomic landscape of lncRNAs identifies xx and xx as novel determinants of xx therapeutic response in rheumatoid arthritis

Genome-wide micro-RNA association study identifies xx and xx as novel determinants of xx therapeutic response in rheumatoid arthritis

Shicheng Guo, Genome-wide DNA methylation identifies novel biomarkers for worse outcome rheumatoid arthritis subtypes