Therapeutic Effects of Infliximab, Tocilizumab and Abatacept in Rheumatoid **Arthritis** OHiroshi IIJIMA¹, Seiji NAKAMURA¹, Katsuya SUZUKI², Yuko HATA¹, Chun Ren LIM¹, Yohei ISHIZAWA¹, Hideto KAMEDA³, Koichi AMANO⁴, Kenichi MATSUBARA¹, Ryo MATOBA¹, Tsutomu TAKEUCHI²

28AB-ISMS05 Identification of Predictive Gene Expression Signatures Associated with the

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While the best outcomes using biologic agents to treat Rheumatoid Arthritis (RA) patients, depends on timely adaptation of therapies, determining first biologic for RA patients still lacks a standard methodology. We studied

the baseline genome-wide gene expression patterns of 209 RA patients' whole blood, prior to administration of infliximab (IFX), tocilizumab (TCZ) or abatacept (ABT). Based on gene set enrichment analysis, we have

identified predictive gene signatures associated with the therapeutic effects of IFX, TCZ and ABT, namely

inflammasome pathway, genes specifically expressed in B cells and in NK cell-specific genes, respectively. We

believe each gene signature is likely related to the target pathway of each biologic agent.