

JCR 2021 ICW Excellent Abstract Award (29名)

Entry No. (演題ID)	Name	Abstract title
20001	Mitsuhiro Akiyama	Effectiveness of tocilizumab in patients with TAFRO syndrome.
20006	Yasuo Nagafuchi	Autoimmunity associated HLAs regulate T cell receptor repertoire in vivo.
20020	Suguru Honda	Polygenic risk scores predict radiographic progression in patients with rheumatoid arthritis
20023	Yasushi Kondo	COVID-19 shares clinical features with anti-melanoma differentiation associated protein 5 positive dermatomyositis and adult Still's disease
20026	Yasuyuki Todoroki	GLUT1-dependent glycolysis in T-bet+CD11c+ B cell differentiation and its role in systemic Lupus erythematosus (SLE)
20037	Aya Nishiwaki	Transcriptome analysis and B cell receptor repertoire in IgG4-Related Disease
20038	Takanori Ichikawa	Interferon-gamma producing immunocompetent cells for predicting disease activity and severity in adult-onset Still's disease
20043	Hideyuki Takahashi	Patient stratification of systemic eosinophilic diseases based on clinical features
20047	Chiharu Hidekawa	Dose hydroxychloroquine have a protective effect on infectious disease in patients with systemic lupus erythematosus or not?: The longitudinal study from the LUNA registry
20050	Takuya Kotani	Examination of the relationship between cerebrovascular disease associated with microscopic polyangiitis and serum cytokine levels
20052	Kuniyuki Aso	Anti-gAChR antibody as a novel biomarker for lupus enteritis in systemic lupus erythematosus
20056	Keita Ninagawa	Quantitative evaluation of chest CT differentiates the dominance of pulmonary vascular disease or interstitial lung disease on hemodynamic abnormalities in systemic sclerosis
20062	Maho Nakazawa	Transcriptome analysis of alveolar macrophages identifies involvement of PD-1/PD-Ls in pathogenesis of rheumatoid arthritis-associated interstitial lung disease
20064	Masaru Yoshimura	Prophylactic effect of salazosulfapyridine for pneumocystis pneumonia in rheumatoid arthritis: A retrospective propensity score-matched cohort study
20066	Naoki Hamada	Report on cases of inflammatory arthritis induced by immune checkpoint inhibitors in our hospital.
20070	Yusuke Miyazaki	Effects of tocilizumab on immunophenotypic features in patients with large vessel vasculitis: results of a 1-year FLOW study
20080	Haruki Matsumoto	Role of T cell immunoglobulin and mucin domain-3 (TIM-3) in determining clinical phenotype of RA
20083	Masataka Umeda	ADAM9 Drives Th17 cell differentiation and Autoimmunity by Activating TGF-beta1
20086	Ayako Yamaguchi	The possibility of drug free remission after sustained remission in patients with RA - The three-year results of the FREE-J study, a real world prospective observational cohort study
20095	Saeko Yamada	Immunomics analysis of rheumatoid arthritis identified pre-dendritic cells as a key cell subset of treatment resistance.
20097	Hiroshi Takei	Contrast of muscle magnetic resonance imaging and pathological findings of muscle tissue in patients with anti-aminoacyl transfer RNA synthetase antibodies
20098	Haruka Takahashi	Analysis of heterogeneity of IL-6 secretion from synovial fibroblasts by cell-to-cell interaction
20109	Shotaro Kawano	TNFR2 signaling enhances suppressive function of Follicular regulatory T cells.
20111	Yuki Kudo	Establishment of iPSc and differentiated endothelial cells of systemic sclerosis associated pulmonary arterial hypertension ; functional and molecular analysis
20112	Akira Onishi	Effectiveness of biological DMARD/JAK inhibitor monotherapy in patients with rheumatoid arthritis - ANSWER longitudinal cohort study -
20117	Yuya Fujita	Approach to establish a novel evaluation method for Neuropsychiatric Systemic Lupus Erythematosus (NPSLE) -LOOPS Registry-
20126	Naoaki Ohkubo	Remission induction therapy with mycophenolate mofetil in patients with lupus nephritis -LOOPS registry-
20128	Tomohiro Koga	Granulocyte-macrophage colony-stimulating factor and tumor necrosis factor alfa in combination is a useful diagnostic biomarker to distinguish familial Mediterranean fever from sepsis
20131	Saori Sakaue	A global atlas of genetic associations of 220 deep phenotypes enabled genetic-driven categorization of autoimmune diseases