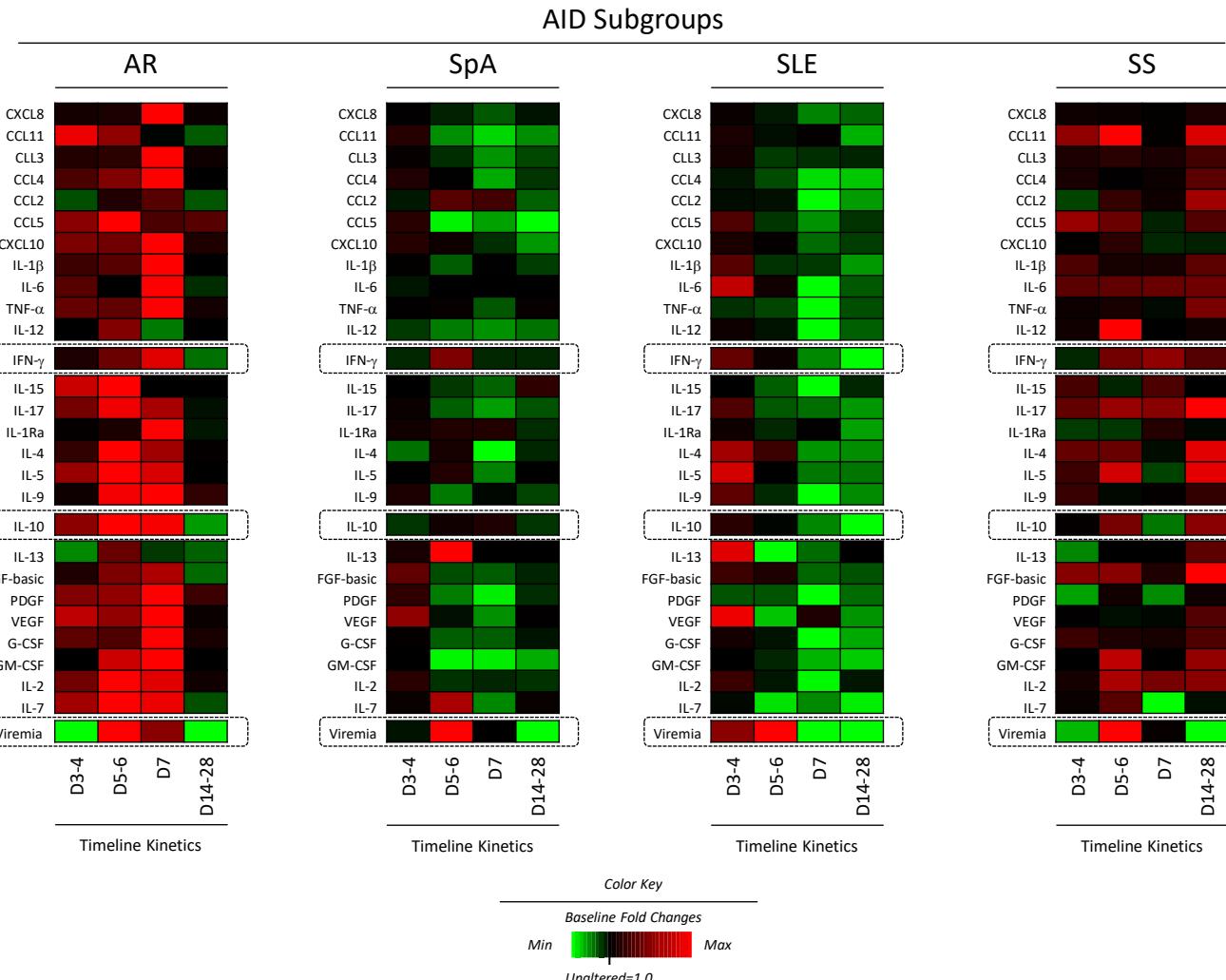


**Serum biomarker profile orchestrating the seroconversion status of patients with
autoimmune diseases upon planned primary 17DD Yellow Fever vaccination**

Ismael Artur da Costa-Rocha^{a*}, Ketty Lysie Libardi Lira Machado^{b*}, Ana Carolina Campi-Azevedo^a, Andréa Teixeira-Carvalho^a, Vanessa Peruhype-Magalhães^a, Sheila Maria Barbosa de Lima^c, Emily Hime Miranda^c, Gisela Freitas Trindade^c, Thays Zanon Casagrande^b, Samira Tatiyama Miyamoto^b, Sávio Carvalho Deotti^b, Rafaela Villa Real Barbosa^b, Priscila Costa Martins Rocha^b, Erica Vieira Serrano^b, Valquiria Garcia Dinis^{b,d}, Sônia Alves Gouvêa^b, Maria Bernadete Renoldi de Oliveira Gavi^b, Lidia Balarini da Silva^b, Ruben Horst Duque^b, Ana Paula Espíndula Gianordoli^b, Maria de Fatima Bissoli^b, Maria da Penha Gomes Gouvea^b, Lauro Ferreira da Silva Pinto-Neto^d, Ana Paula Neves Burian^e, Francieli Fontana Sutile Tardetti Fantinato^f, Gecilmara Salviato Pileggi^g, Licia Maria Henrique da Mota^h, Valéria Valim^{b#} and Olindo Assis Martins-Filho^{a#}

Heatmap Timeline Kinetic Profiles of Serum Biomarkers After 17DD-YF Primary Vaccination
According to the Type of Autoimmune Disease



Supplementary Figure 1 - Heatmap Timeline Kinetic Profiles of Serum Biomarkers After 17DD-YF Primary Vaccination According to the Type of Autoimmune Disease. Heatmaps were constructed considering the baseline fold change values at each time point along the kinetic follow-up (D3-4, D5-6, D7 and D14-28). This approach was employed to draw the overall change in the serum biomarkers profile after primary 17DD-YF vaccination of Autoimmune Disease patients segregated according to the type of autoimmune disease [Rheumatoid Arthritis (RA, n=38), Spondyloarthritis (SpA, n=51), Systemic Lupus Erythematosus (SLE, n=21) and Sjögren's Syndrome (SS, n=30)]. Data interpretation was carried out based on the color keys were employed underscore the baseline fold value = 1.0 as the reference for unaltered levels (■), the baseline fold value <1.0 for decreased levels (■) and the baseline fold value >1.0 for increased levels (■), according to the paired sample collected at D0.