

Figure S1. Uncut gels. Representative western blot for tissue factor and ferritin of serum samples of the subjects obtained from COVID-19 patients and HC. The samples were run blindly. The graphs in the Figures 3-5 (position C) were obtained after the densitometry analysis of the western blot. In blue: discharged patients; in red: deceased patients; in green: healthy controls; and in black: patients without medical records and excluded to the cohort.

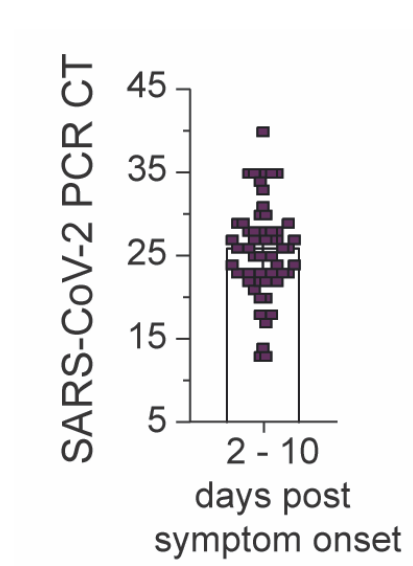


Figure S2. SARS-CoV-2 PCR cycle threshold (CT) of nasopharyngeal swab after symptom onset of patients' admission in MDNH.

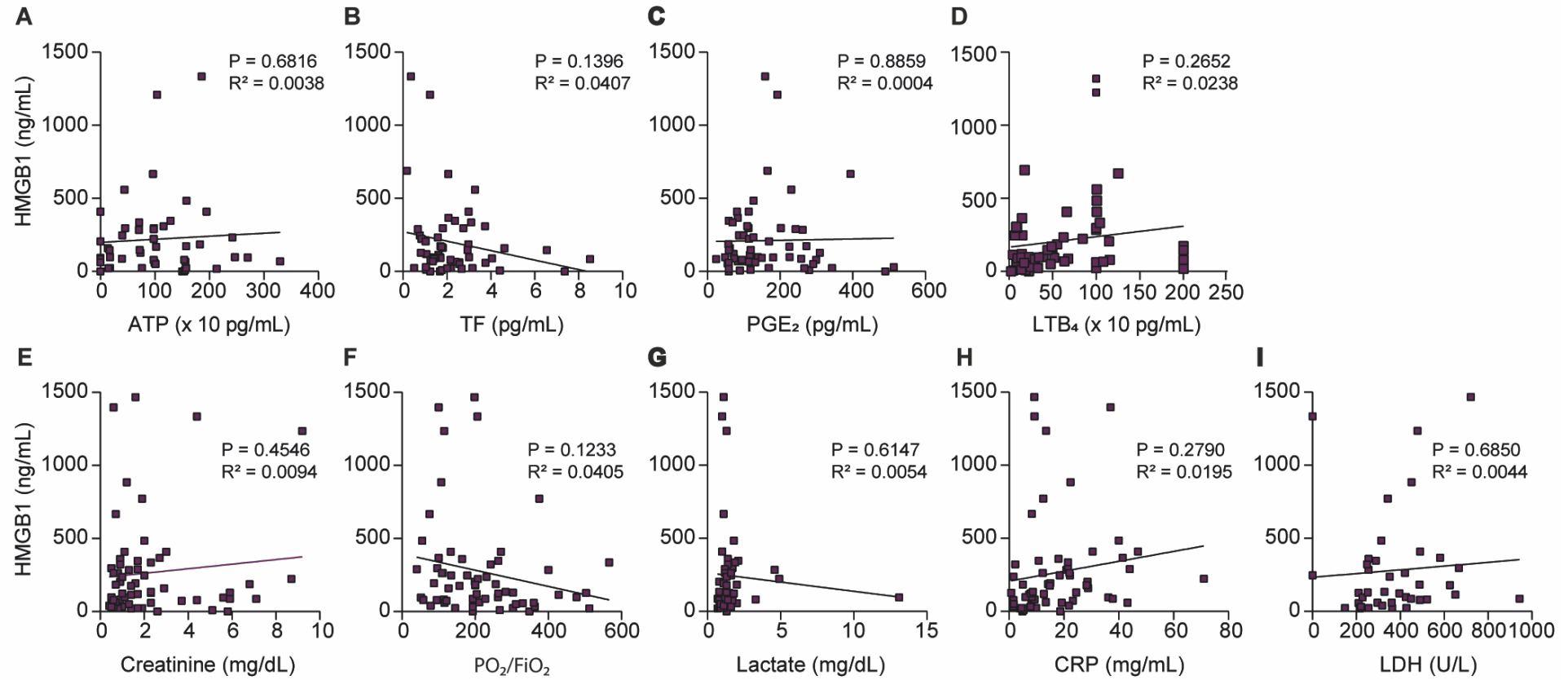


Figure S3. Spearman's correlation analyses between inflammatory mediators and tissue injury biomarkers with HMGB1. (A-I) Spearman's correlation analyses between HMGB1 serum levels versus ATP, TF, PGE₂, LTB₄, creatinine, PO₂/FiO₂, lactate, CRP, and LDH levels. We plotted the highest value obtained for HMGB1 after analysis of two or three samples received from each COVID-19 patient. The values for ATP, TF, PGE₂, LTB₄, creatinine, PO₂/FiO₂, lactate, CRP, and LDH were paired exactly with the same sample chosen for analysis of HMGB1. No correlations were detected with the analyzed parameters.