

Comparison of Human Serum and Plasma by Two-Dimensional Gel Electrophoresis

Hyun-Jung Kim, Eun-Jung So, Ji-Su Kim, Mi-Ryung Kim and Chan-Wha Kim

School of Life Sciences and Biotechnology, Korea University, Seoul 136-701, Korea.

TEL: +82-2-3290-3439, FAX: +82-2-3290-3957, E-mail: cwkim@korea.ac.kr

Serum or plasma proteins are the indicators of patient's physiological or clinical status. Proteins involved in coagulation factors present in high abundance in serum or plasma, have been functionally characterized and associated with various disease states¹⁾. Serum is the blood plasma without fibrinogen and other clotting factors. Plasma is the fluid portion obtained after centrifugation of the anticoagulated specimen. Generally it is expected that there are differences in the protein profile between serum proteins and anti-coagulant matter (ACM; EDTA, heparin or trisodium citrate) treated-plasma proteins. In this study, we removed the most abundant serum proteins, albumin and IgG using a removal kit prior to display human blood serum or plasma proteins with two-dimensional electrophoresis (2-DE) gels. The proteins which show the differences in spot intensities in 2-dimensional electrophoresis gel were identified with ESI-Q-TOF MS/MS. Our study will provide the good guidance for selecting either serum or plasma samples for clinical and biomedical research.

Reference

1. Pieper R Gatlin CL Makusky AJ Russo PS Schatz CR Miller SS Su Q McGrath AM Estock MA Parmar PP Zhao M Huang ST Zhou J Wang F Esquer-Blasco R Anderson NL Taylor J Steiner S. Rembert pieper Christine LMozdarani H Kamali S. The human serum proteome: display of nearly 3700 chromatographically separated protein spots on two-dimensional electrophoresis gels and identification of 325 distinct proteins (2003), *Proteomics*, 3(7), 1345-1364.