Supplementary Material

Unlocking Chromatographic Quantification of Serum Lactate Dehydrogenase via Ionic-Liquid-Based Aqueous Biphasic System Pretreatment

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Tables

Table S1. Numerical data for depletion efficiency of high-abundance serum proteins (DE_{PROT}%), extraction efficiency of LDH (EE_{LDH}%), recovery yield of LDH (RY_{LDH}%) and relative error (RE%) across different compositions of the initial mixtures used to form ABS. Values are reported as mean ± standard deviation (σ) of independent replicates. In cases where all replicates yielded identical values, the standard deviation is not shown.

IL	Weight percentage composition / (wt %)		$DE\%_{Prot} \pm \sigma$	$EE\%_{LDH} \pm \sigma$	$RY\%_{LDH} \pm \sigma$	$RE\% \pm \sigma$
	IL	$K_3C_6H_5O_7$/ $C_6H_8O_7$ buffer solution				
$[P_{4444}]Cl$	30	60	33.10±0.06	100	100	-
$[P_{4444}]Br$	30	60	100	100	82 ± 2	18 ± 2
	30	20	100	100	40 ± 1	60 ± 1
	30	40	100	100	65.5 ± 0.3	34.5 ± 0.3
	45	40	100	100	92 ± 2	8 ± 2
	15	40	100	100	32.3 ± 0.7	67.6 ± 0.7

Figures

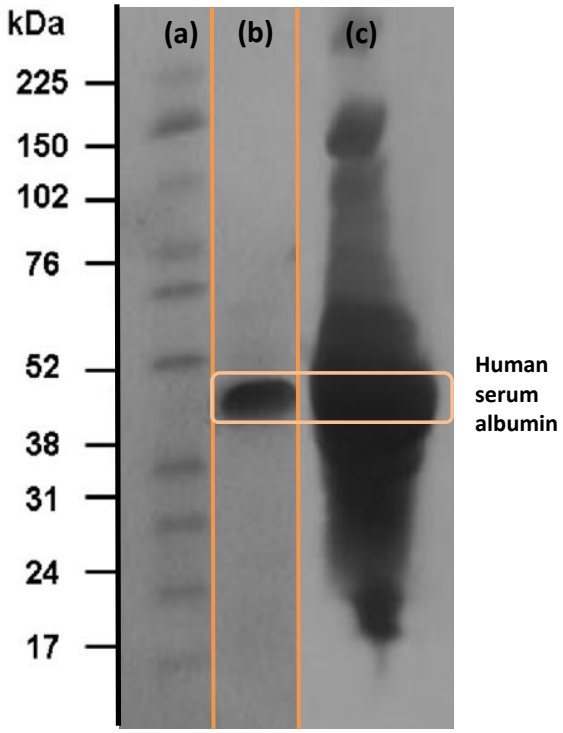

Fig. S1. SDS-PAGE analysis of serum proteins: (a) molecular weight marker; (b) HSA sample, 1g.L^{-1}; (c) human serum proteins precipitated at interphase of the IL-based ABS (diluted 2:1 in PBS buffer).