



Figure S2. SSL3 peptide response curves. Dilution curve of SSL3 peptides spiked in human plasma backgrounds (1 μ g on column) at concentrations ranging from ~100 fmol to ~5 amol SSL3 on column and analyzed with SRM-MS targeting peptides previously detected with shotgun MS. The peptide total area is calculated by summing the area-under-the curve of all transitions per precursor ion. The plus signs (+) indicates peptide charge state and numbers within brackets ([]) indicate mass change (modification) of the preceding amino acid residue. The three selected peptide sequences are shown in *bold*.