The removal of Triton X-100 by dialysis is feasible!

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Fig. S1 Comparison between positive ion MALDI-TOF mass spectra of trypsin digested carbonic anhydrase (CA) without TX (a), after addition of 0.1% TX (b) and after addition of 0.5% TX and subsequent removal of the detergent by dialysis for 72 h. Red dots indicate all CA matching peptide masses.
Fig. S2 Comparison between positive ion MALDI-TOF mass spectra of trypsin digested bovine serum albumin (BSA) without TX (a), after addition of 0.1% TX (b) and after addition of 0.5% TX and subsequent removal of the detergent by dialysis for 72 h. Red dots indicate all BSA matching peptide masses.
Fig. S3 Effect of increasing concentrations of TX on positive ion MALDI-TOF mass spectra of a trypsin digested protein fraction of blood plasma (sample 1)
**Fig. S4** Positive ion MALDI-TOF mass spectra of a trypsin digested protein fraction of blood plasma (sample 2)
Aliquots of protein fraction were prepared without (a) and with addition of 0.5% of TX. TX was removed either by microdialysis for 72h in presence of 5M urea before trypsin digestion (b) or by ethyl acetate extraction after trypsin digestion (c)