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Electronic Supplementary Information

ACCURATE QUANTIFICATION OF CARBOPLATIN ADDUCTS WITH SERUM PROTEINS BY MONOLITHIC CHROMATOGRAPHY COUPLED TO ICPMS WITH ISOTOPE DILUTION ANALYSIS

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Figure ESI-1. Fractionation of whole blood incubated with cisplatin

Figure ESI-2. Workflow for calculation of reaction yields of incubation of proteins standards with carboplatin using different approaches (A and B).

Figure ESI-3. Preparation and characterization of ^{nat}carboplatin-HSA calibrant and ^{194}carboplatin-HSA spike

Figure ESI-4. Incubation of ERM®- DA470k/IFCC serum with carboplatin and Double Species-specific IDA HPLC-ICPMS for quantification of Pt-HSA adducts in the model sample

Figure ESI-1. Fractionation of whole blood incubated with cisplatin

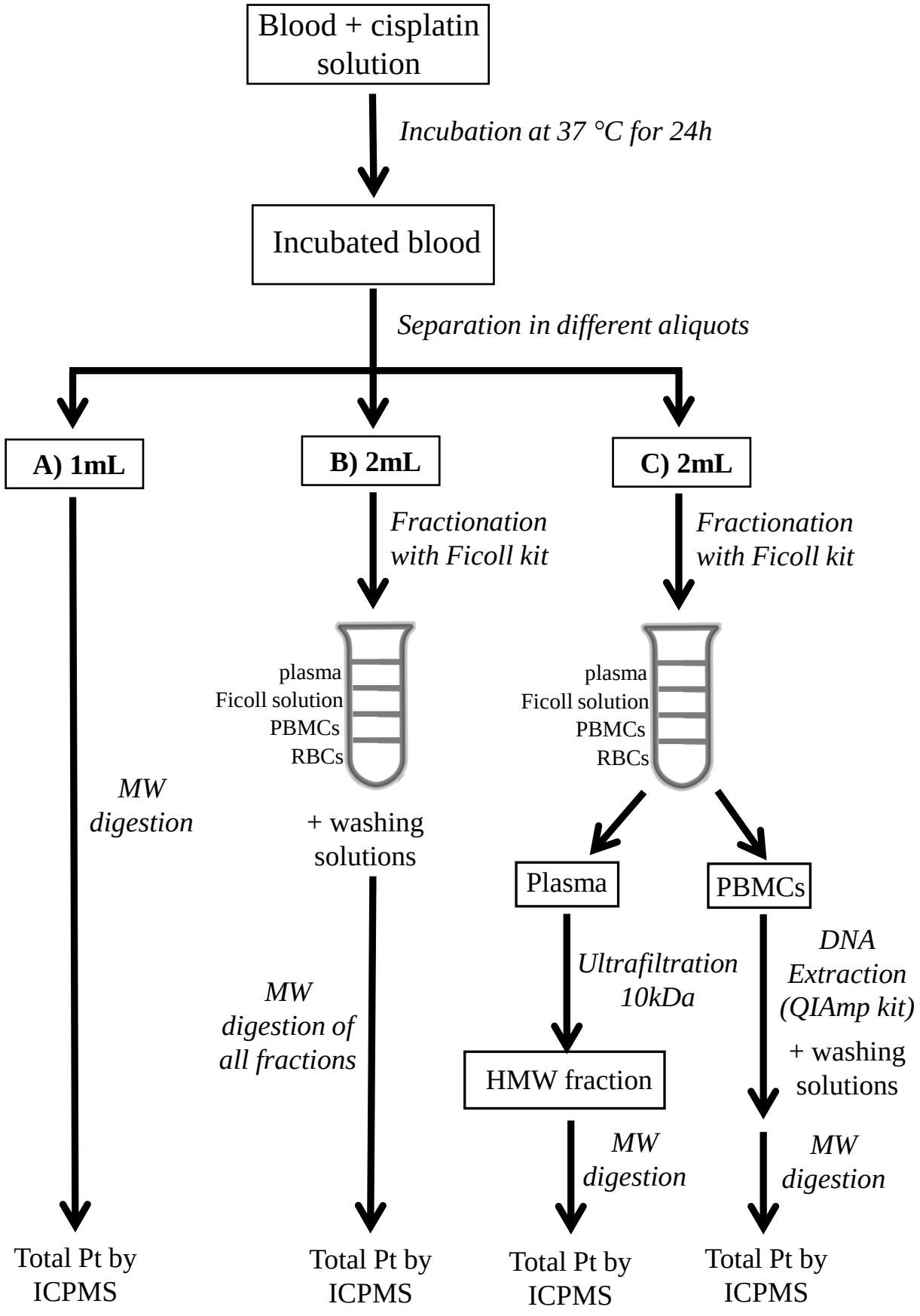
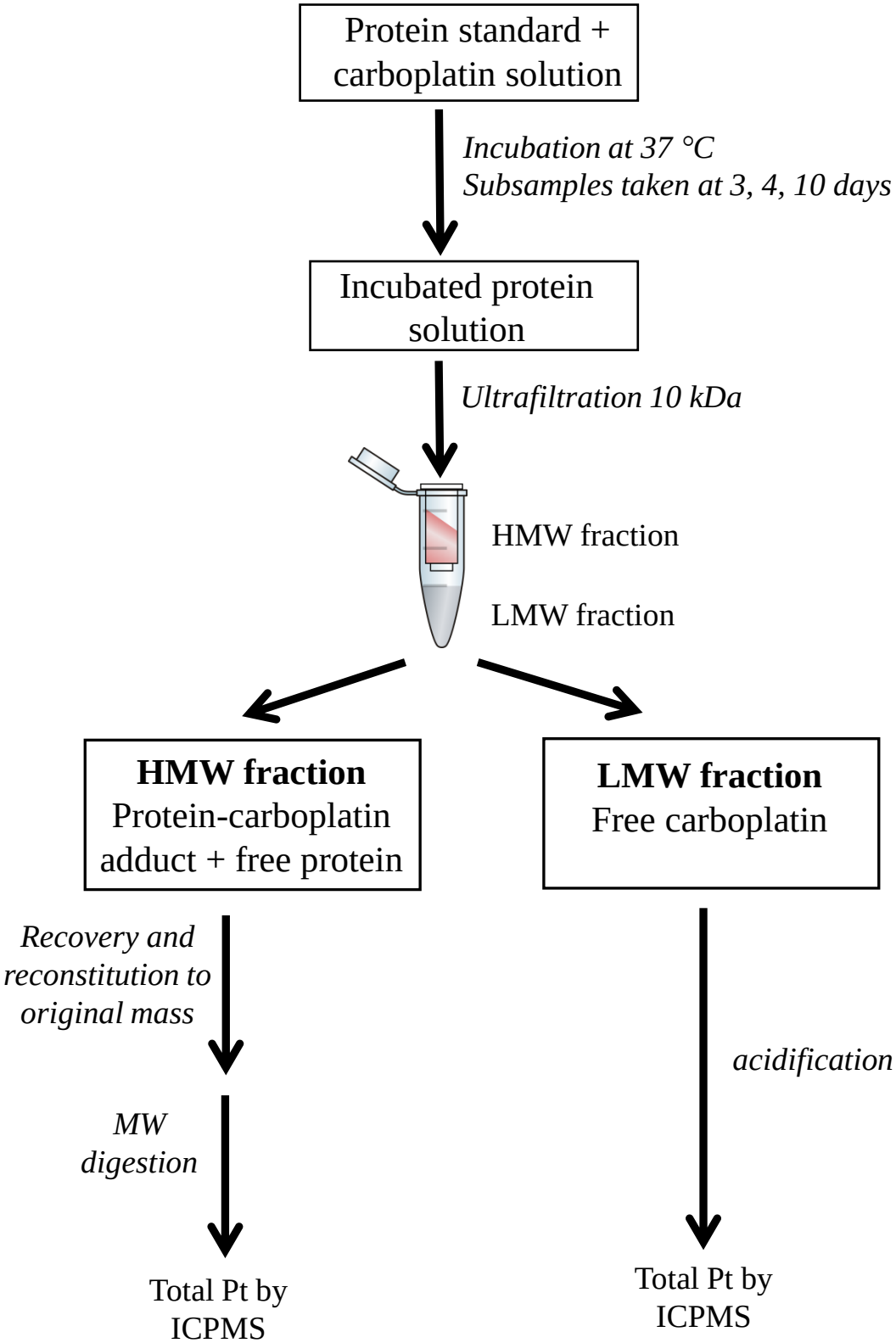


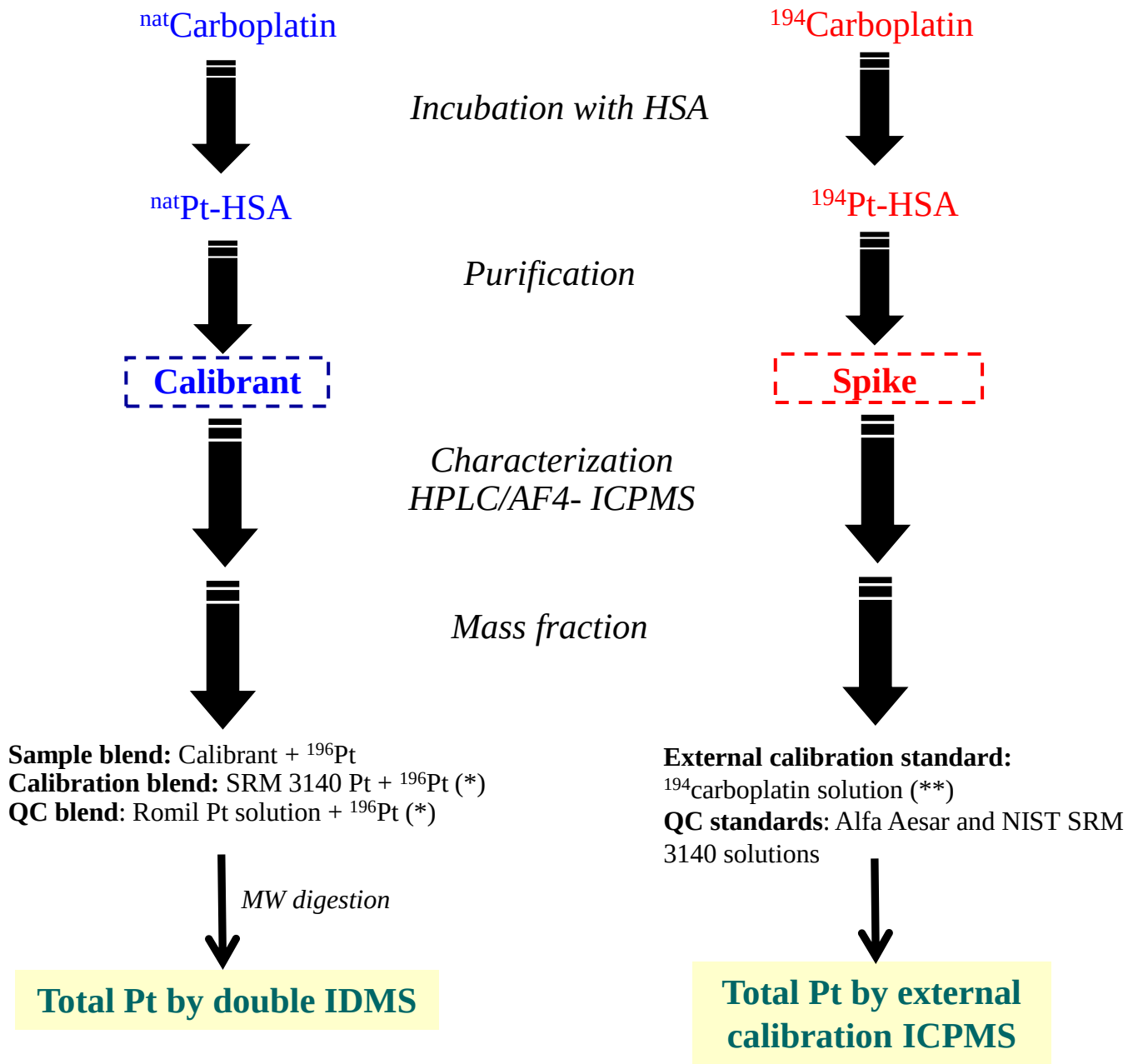
Figure ESI-2. Workflow for calculation of reaction yields of incubation of proteins standards with carboplatin using different approaches (A and B).



A) Yield of adducts formation = Pt conc in HMW*100/ total Pt in incubation solution

B) Yield of adducts formation = 100 - (free carboplatin Pt in LMW*100/ total Pt in incubation solution)

Figure ESI-3. Preparation and characterization of ^{nat}carboplatin-HSA calibrant and ^{194}carboplatin-HSA spike



(*) Calibration and QC blends were analyzed undigested to calculate recovery of MW digestion process

(**) ^{194}carboplatin solution previously characterized in total Pt content by external calibration ICP-OES

Figure ESI-4. Incubation of ERM®- DA470k/IFCC serum with carboplatin and Double Species-specific IDA HPLC-ICPMS for quantification of Pt-HSA adducts in the model sample