





Canine/feline serum or plasma deproteinization for amino acid analysis on Biochrom

COMMENTS 0

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**ABSTRACT** 

This protocol describes the deproteinization process that all canine and feline serum and plasma research samples undergo prior to amino acid analysis with a Biochrom 30+ Amino Acid Analyzer at the Gastrointestinal Lab, Texas A&M University. L-norleucine is used as an internal standard.

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**KEYWORDS** 

amino acids, deproteinization

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MATERIALS TEXT

## **MATERIALS**

- 🔀 750µL Nonsterile Micro-Centrifugal PVDF Membrane Filters, 0.2µm pore size Thermo Fisher Catalog #F2517-5 Step 5
- Amino acid standards physiological acidics and neutrals Millipore Sigma Catalog #A6407
- X Amino acid standards physiological basics Millipore Sigma Catalog #A6282
- **⊠** 5-Sulfosalicylic acid dihydrate BioXtra >=99.0% **Sigma − Aldrich Catalog #S7422** Step 2

- 1 Aliquot 250 µl serum or plasma into 1.5 ml microcentrifuge tube.
- 2 Add 250 μl [5% Sulfosalicylic acid (w/v), 500 μM L-Norleucine] to 1.5 ml microcentrifuge tube. Note: if less than 250 μl serum or plasma is available, can use as little as 100 μl and add reagent in 1:1 (v/v) ratio. i.e., if using 150 μl serum, would add 150 μl [5% Sulfosalicylic acid (w/v), 500 μM L-Norleucine].
- 2.1 To make SSA solution, weigh 5 g sulfosalicylic acid into graduated cylinder, add 6.6 +/- 0.1 mg L-Norleucine, then fill to 100 mL with Biochrom lithium loading buffer. Stir briefly on magnetic stir plate until all solids are fully dissolved. Store at 4°C.
- 3 Vortex each on high speed for 5-10 seconds, and then store in 4°C for 10 minutes.
- 4 Centrifuge 1.5 ml tubes at 10,000 rcf, 4°C for 5 minutes. (2) 10000 x g

**Expected result** 

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5	Transfer supernatant to centrifuge filter tubes, being careful not to disturb pellet or floating lipid layer.
	750μL Nonsterile Micro-Centrifugal PVDF Membrane Filters, 0.2μm pore size <b>Thermo Fisher Catalog #F2517-5</b>

6 Centrifuge filter tubes at 10,000 rcf, 4°C for 5 minutes. 10000 x g Discard filter and store filtrate at -80°C until analysis.

Expected result