

# Struvite Stones and the Microbiome

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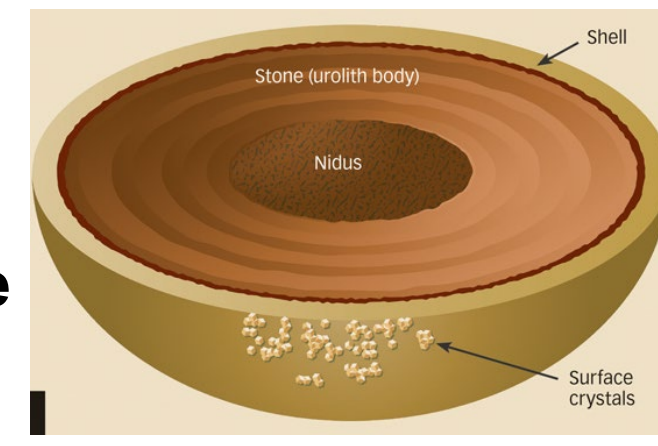
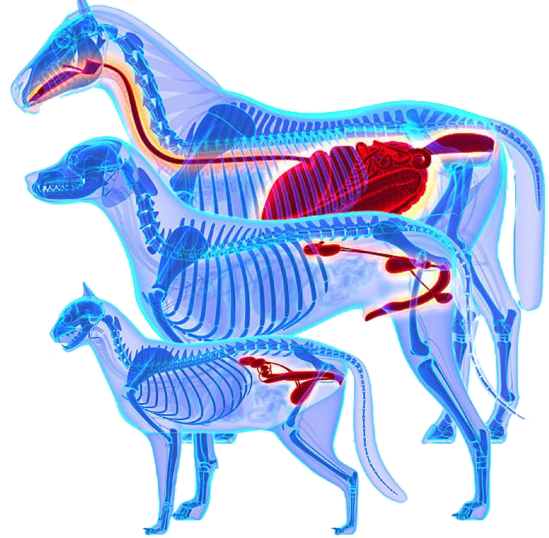
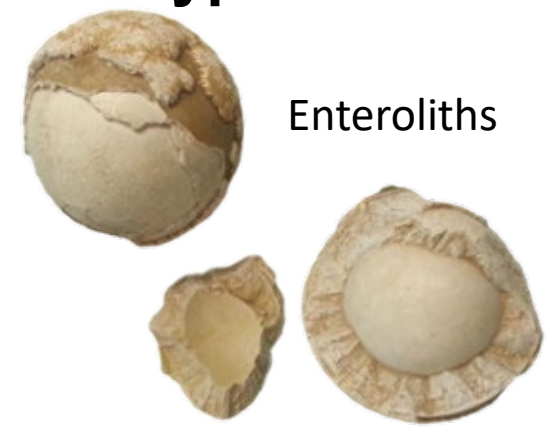
## Introduction

### Enteroliths in Horses and Struvite Uroliths in Dogs and Cats

Endogenously produced mineral concretions, form in concentric rings around a foreign object (nidus). Enteroliths occur in the right dorsal colon of horses while uroliths occur along the urinary tract of dogs and cats.

Primary component= Struvite

- A composite of magnesium ammonium phosphate hexahydrate  $[(NH_4)MgPO_4 \cdot 6H_2O]$
- Alkaline pH is favorable for precipitation
- Struvite uroliths are the most common type of urinary stone in dogs, 2<sup>nd</sup> most common type in cats.



In dogs, virtually all struvite uroliths are infection induced.

Most common types of bacteria:

- Staphylococcus pseudintermedius*
- Proteus mirabilis*
- Klebsiella* spp. (known N fixer)

Two ways to produce  $NH_3$ :

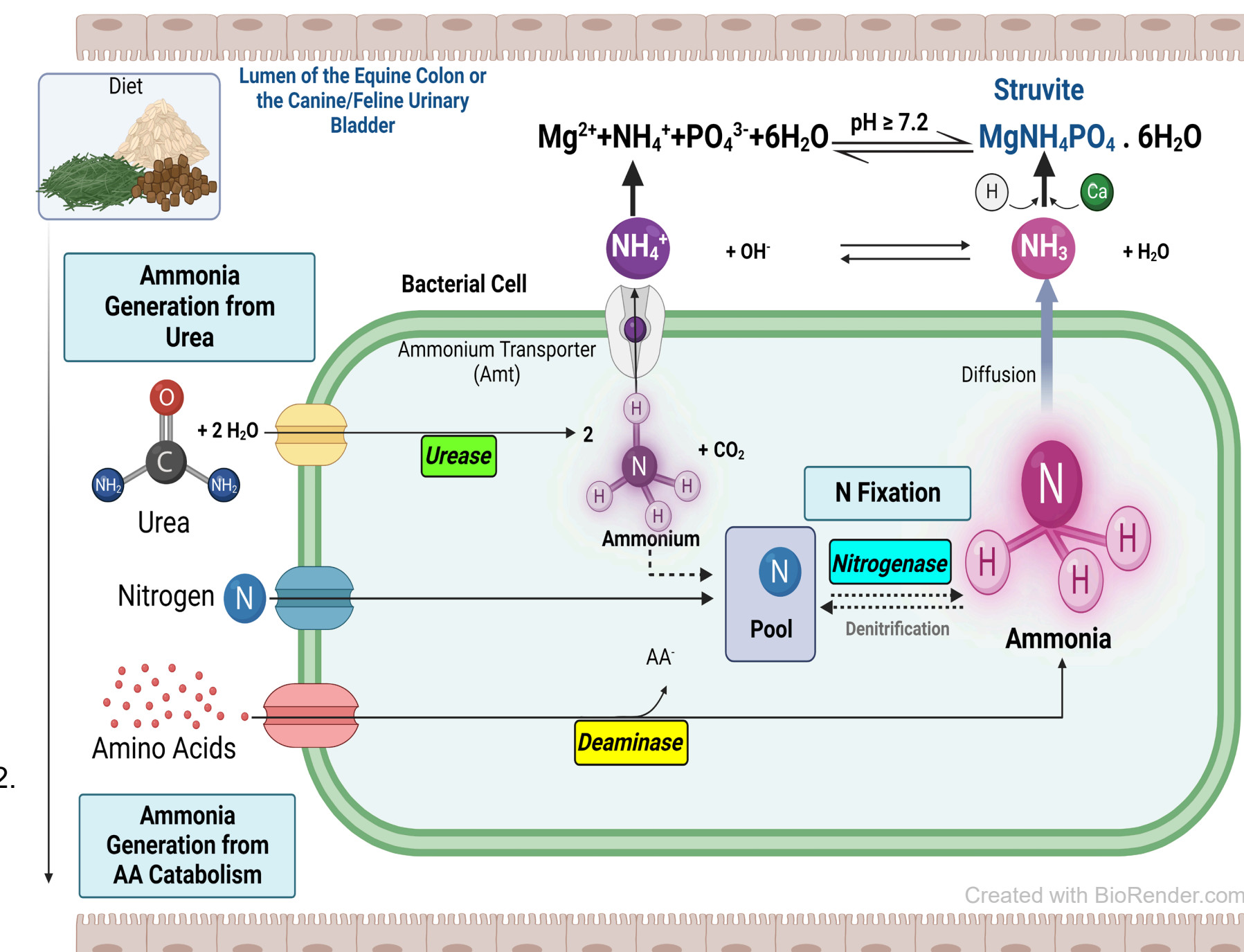
Urease producers- *Staph.* and *Proteus*

Associated with UTIs and struvite formation

Urease  $\rightarrow$  hydrolysis of urea, forming  $NH_3$  and  $CO_2$ .

N Fixers-*Klebsiella*

Nitrogenase  $\rightarrow$  microbial production of  $NH_3$



## Purpose

- Urolithiasis occurs in a vast array of species
- Primary stones that occur in dogs and cats are struvite and CaOx
- Struvite is the common denominator, but stone characteristics have not been compared at depth
- Insight into the disease processes can be used in the clinical setting, optimizing treatment options
- Methods can be applied to other types of urinary stones.
- Translational applications exist for both humans and many veterinary species.

## Objectives

Hypotheses:

- A microbial community exists (microbiome) within struvite urinary stones in both dogs and cats that contributes to stone formation.
- Based on previous findings, we postulate that a similar microbiome occurs in equine enteroliths and dog/cat struvite urinary stones. Members of the microbiomes contribute to an environment that favors struvite precipitation and stone formation.

Objective: Identify the microbiome of canine/feline struvite urinary stones via deep RNA metagenomic sequencing.

## Methods

### Sample Selection

Samples selected using the stone database of the Gerald V. Ling Urinary Stone Analysis Laboratory at UC Davis. For each sample, the mineral composition of the layers and the core had been determined by the lab using optical crystallography with polarizing light microscopy.

Mineral composition was displayed as seen here:

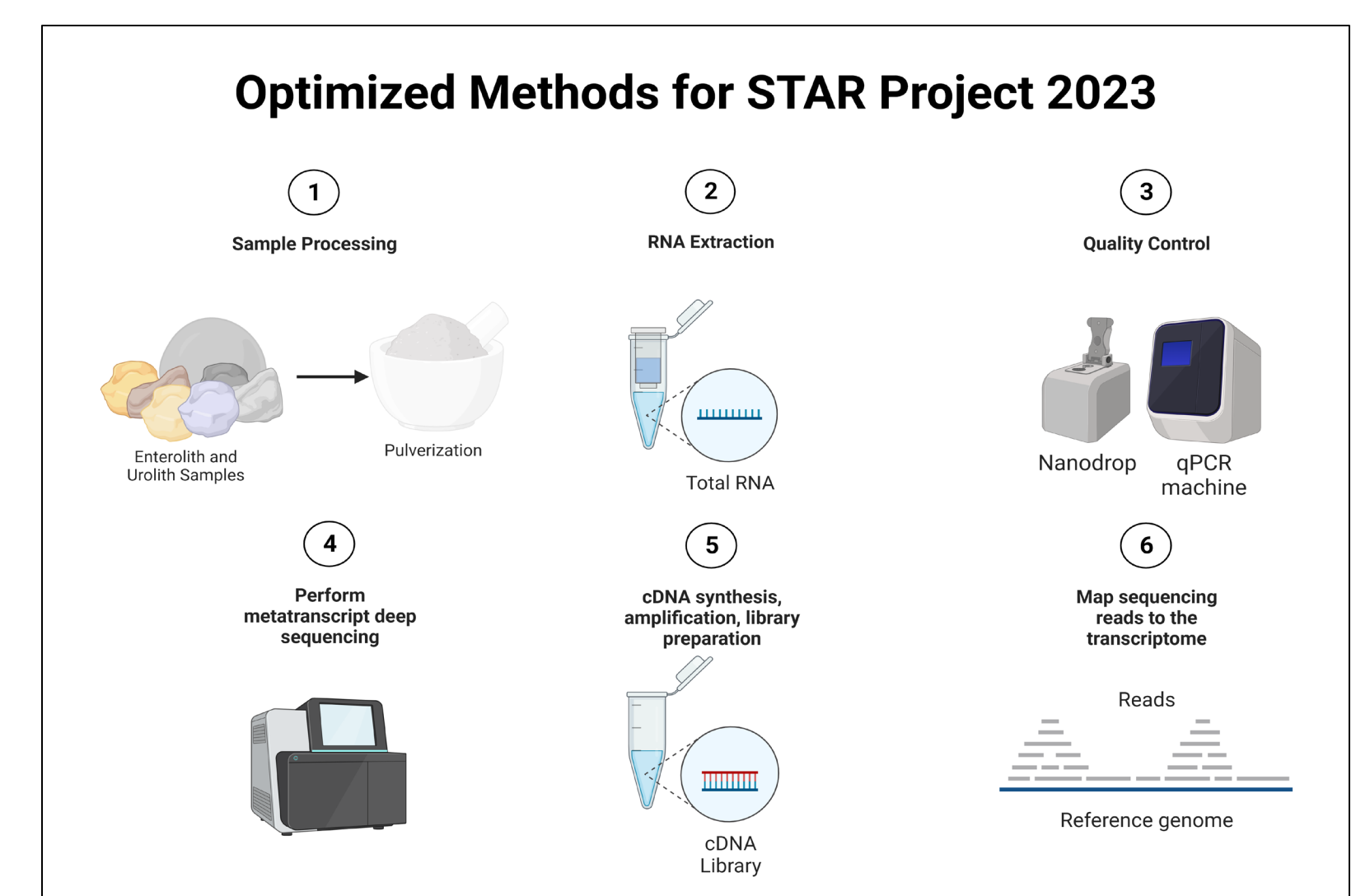
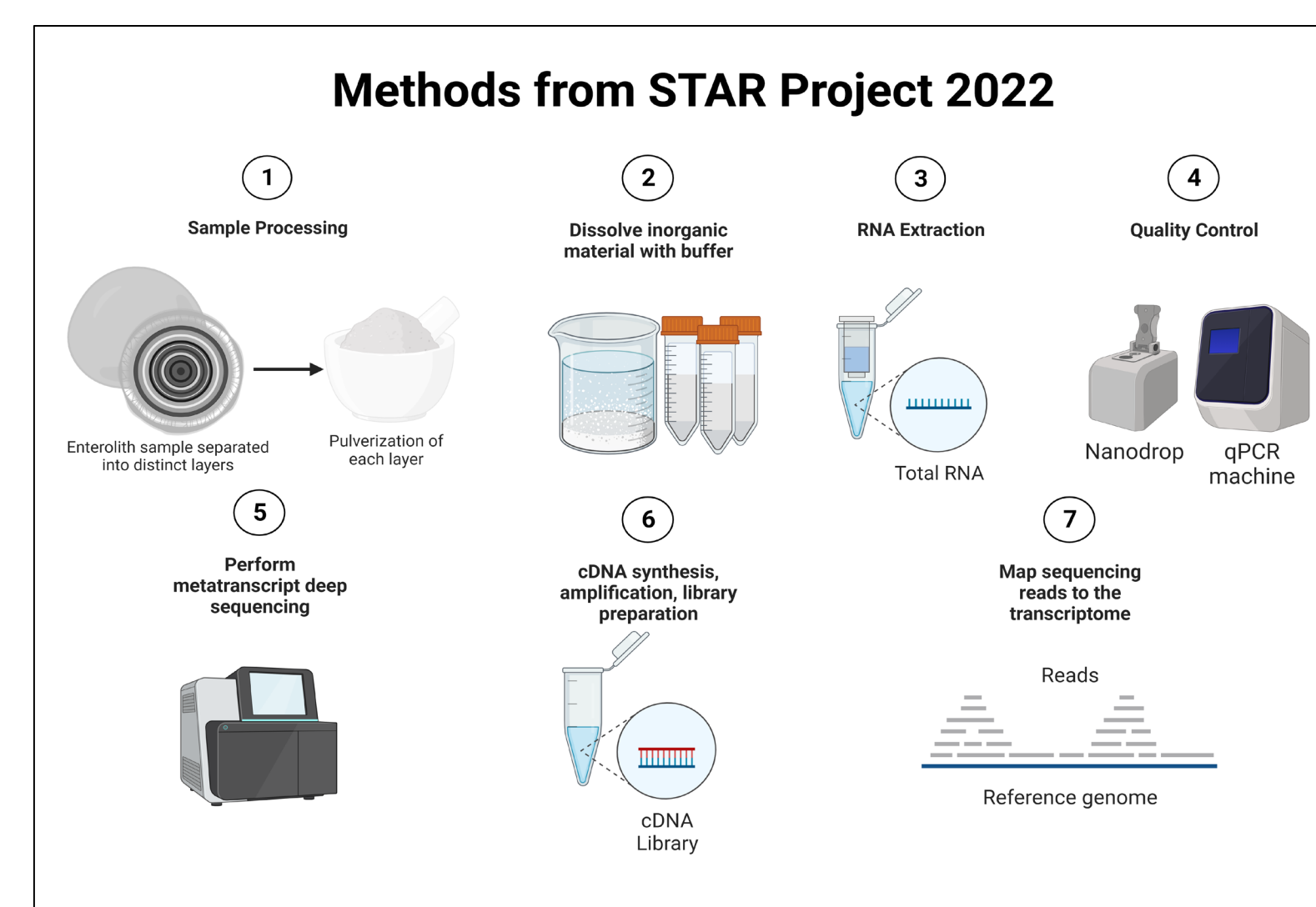
SPECIES	COMP. OUTR	LAYER_1	LAYER_2	LAYER_3	CORE
CANINE	struvite 100%	struvite 95%, apatite 5%	struvite 99%, apatite 1%	-	struvite 100%

Samples were selected from the lab's database based on the following criteria:

- Species (canine, feline, or equine)
- Urine culture performed
- Stone culture performed
- Layers and the core composed primarily of struvite (some carbonate apatite was acceptable)



## Results: RNA Extraction Optimization



By optimizing the methods used for RNA extraction we were able to increase both the concentration and the quality of the RNA extracted.

- Eliminating step 2 preserves more sample RNA.
- The acidic buffer likely degraded some of the RNA previously.
- Pulverizing the stones was effective for quality RNA extraction.
- Each sample was crushed whole to detect the presence or absence of RNA within each stone.
- Separating different urolith layers by hand is not feasible.

## Results: RNA Extraction from Enteroliths and Uroliths

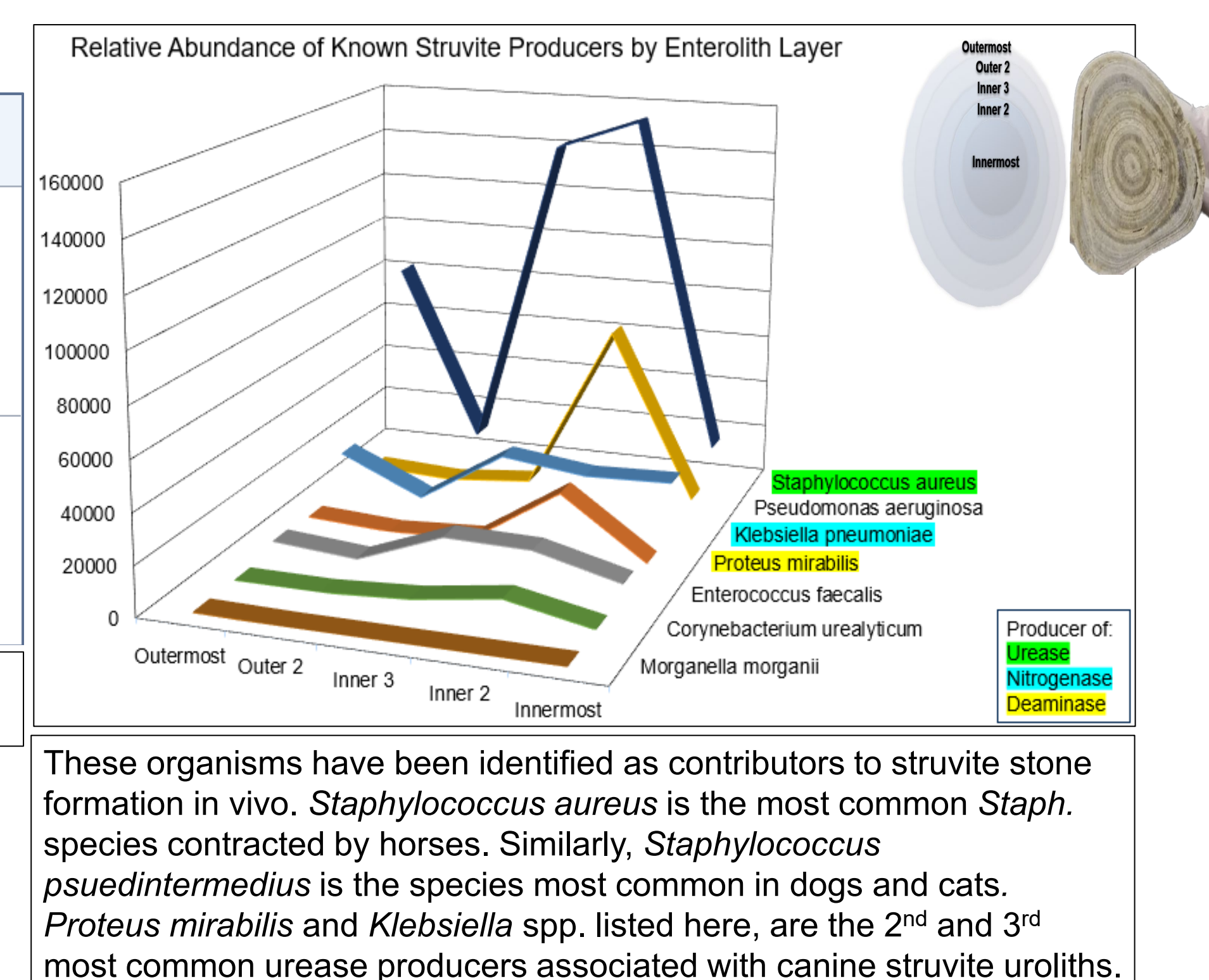
### Nanodrop Results by Species

	RNA Conc. (ng/μl)	Protein (260/280)	Organics (260/230)
Canine Urolith	103.5	1.98	2.19
Feline Urolith	201.0	1.85	2.04
Equine Enterolith	54.4	1.62	0.50

### Comparison of Enterolith RNA Extraction Methods

Enterolith Sample	RNA Conc. (ng/μl)	Protein (260/280)	Organics (260/230)
Processed with buffer (2022)	27.7	1.50	0.10
Processed without buffer (2023)	54.4	1.62	0.50

The optimized extraction methods were found to produce an RNA sample of higher concentration and quality for deep sequencing.



## Conclusion and Future Directions

Future Directions :

Deep RNA metagenomic sequencing is currently underway for the canine and feline urolith samples along with more enterolith samples.

- Determine the relative abundance of struvite producers in both the canine and feline uroliths and compare to enterolith data.
- Work to determine function using transcripts from the metagenome. The transcripts allow identification of genes associated urease, but also broader  $NH_3$  and  $CO_2$  metabolism associated with struvite formation.
- Apply these techniques to other types of uroliths

Conclusions:

- An active microbiome is present in enteroliths and canine/feline struvite uroliths
- Enteroliths and struvite uroliths contain quality, extractable RNA
- The sample preparation is a key aspect of analyzing these sample types
- The stones contain a microbial community that likely helps facilitate struvite formation both in the equine GI tract and the canine/feline urinary systems

## Acknowledgements

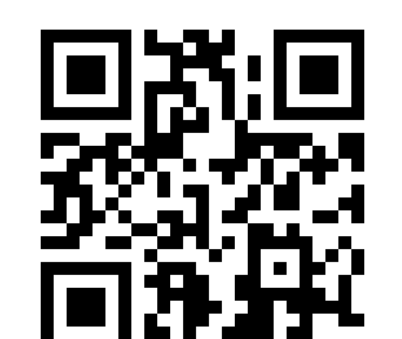
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