Isolation and Characterization of Three Bacteriophages Targeting Acinetobacter baumannii AB5075 **Tenly Hansen, M.S.B.S. '24** Carolyn Schroeder, M.S. Michael Carruthers, Ph.D. Department of Microbiology and Immunology | College of Osteopathic Medicine

Background

- Acinetobacter baumannii
- Responsible for 90% of all reported *Acinetobacter* infections in humans¹
- Primarily associated with nosocomial infections in immunocompromised patients
- Accounts for 20% of infections across ICUs worldwide, with a mortality rate greater than 50%²
- Common clinical manifestations:
- Urinary tract, respiratory tract, gastrointestinal tract, wound, and bloodstream infections³
- Carbapenem-resistant *A. baumannii is* classified as a critical priority pathogen^{2,}
- Pan-drug resistant strains have been isolated

Bacteriophages

- Viruses that infect bacteria and archaea
- Most abundant organisms on Earth (10³¹)

Phage Therapy

- Use of lytic bacteriophages to selectively target and kill pathogenic bacteria causing infection
- Promising advantages include:
- High host specificity⁴, low toxicity⁴, auto-dosing⁵, and biofilm disruption⁶
- Only 184 phages targeting *A. baumannii* with publicly available genome sequences

Phage Morphological Assessment



Isolation of Novel Phages



or super translucent morphotypes to assess plaquing.

200nm

Figure 2: Representative TEM micrographs of phages DMU2, DMU3, and DMU4 exhibit myophage morphologies (i.e., icosahedral heads and contractile tails).

Figure 1A: Isolation of Bacteriophages from Activated Sludge: obtained at the Des Moines Metropolitan Wastewater Reclamation Authority. Figure 1B: Spot Titrations of Acinetobacter Phages: DMU2, DMU3, and DMU4 on soft agar overlays containing A. baumannii AB5075 opaque

200nm



Conclusions

- Isolated three novel phages that target the model *Acinetobacter baumannii* strain, AB5075
- Each phage exhibits a myophage morphology (icosahedral head, contractile tails) with unique dimensions (data not shown)
- This work advances our research into phages that target A. baumannii, adding to a growing collection of phages for potential therapeutic use against this notoriously antibiotic-resistant pathogen

Future Work

- Determine phage host range using a diverse collection of 100 *A. baumannii* clinical isolates
- Cultured from 2,533 patients across 4 continents by the MRSN from 2001-2017; includes multi- and pan-drug resistant isolates
- Sequence and characterize the genomes of phages
- Facilitates screening phage genomes for virulence factors, antibiotic resistance genes, and DNA recombination machinery
- Determine phage growth kinetics such as adsorption rate, latency period, and burst size

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200nm