





Table 1. Characteristics of Island3 and Sbash Genomes

Phage	Genome Size (bp)	Cluster	% GC	Number of Genes	Isolation Location	Isolation Institution
Island3	47287	11	66.8	76	Oakland, PA	Carnegie Mellon University
Sbash	55832	12	65.6	89	Durban, South Africa	University of Kwazulu-Natal

Table 2. Clusters and hosts of phages used in this study.

Phage	Abbreviation used in	Cluster	Host
Antonia	A	B1	Mycobacterium smegmatis
Beelzebub	Bz	S	Mycobacterium smegmatis
Butters	В	Ν	Mycobacterium smegmatis
CBorch11	СВ	H1	Mycobacterium smegmatis
Deadpool	D	Unclustered	Microbacterium foliorum
DrLupo	DL	H2	Mycobacterium smegmatis
Giles	G	Q	Mycobacterium smegmatis
Island3	13	11	Mycobacterium smegmatis
JacoRen57	JR	AB	Mycobacterium smegmatis
Kebs	Ке	Unclustered	Gordonia terrae
Knocker	К	B9	Mycobacterium smegmatis
Larva	L	К5	Mycobacterium smegmatis
Quesadilla	Q	В9	Mycobacterium smegmatis
RedRaider77	RR	S	Mycobacterium smegmatis
Roots515	R	C1	Mycobacterium smegmatis
Sbash	Sb	12	Mycobacterium smegmatis
Sibs6	S	A1	Mycobacterium smegmatis
Smoothie	Sm	Unclustered	Microbacterium foliorum
Tortellini	Т	P2	Mycobacterium smegmatis
Wrigley	W	CY	Gordonia terrae

Introduction

With antibiotic resistance on the rise, phage therapy presents a new approach to fighting bacterial infections. Phages are viruses that exclusively infect bacteria. When a lytic phage infects a bacterium, the result is lysis of the host bacterial cell; this makes them useful in a clinical setting. Through programs such as SEA Discovery, new phages are constantly being discovered, annotated, and archived

SBash and Island3 are cluster I phages that infect Mycobacterium smegmatis and share considerable genetic identity. SDS-PAGE gels were run to isolate proteins from these phages and determine their molecular weight. We measured protein concentrations with a protein assay and injected the isolated proteins into female BALB/c mice. Another group of mice was immunized with phage high titer lysate (HTL). We collected blood from the tail veins of immunized mice, allowed the blood to clot, then centrifuged it to get serum. We performed ELISA assays to determine if phage and protein-specific antibodies were present in the serum and Western blot to investigate how Island3 and Sbash compare from a biochemical perspective. Learning more about the biochemical relationships between phages is important for advancing the field of phage therapy.

Materials and Methods







-teller et al. 2022. G3 (Bethesda). 2022 Jul 29;12(8):jkac140. doi: 10.1093/g3journal/jkac140. PMID: 35727726; PMCID: PMC9339283 Issell and Hatfull. 2017. Bioinformatics 33:784–786. rathdee, et al. 2023. Cell 186: 17ded.). (2014). Cold Spring Harbor, NY: Cold Spring Harbor Lab. Press

atfull, G. F. (2015). Dark Matter of the Biosphere: the amazing world of bacteriophage diversity. J Virol 89:8107-8110 he nature of mycobacteriophage diversity and host preference. Virology 434:187-201. avrich, T. N., & Hatfull, G. F. (2017). Bacteriophage evolution differs by host, lifestyle and genome. Nature Microbiology, 2, 17112.

edulla, et al., (2003). Origins of highly mosaic mycobacteriophage genomes. Cell, 113:171-782. Pope, et al., (2015). Whole genome comparison of a large collection of mycobacteriophages reveals a continuum of phage genetic diversity eLife 4:e064

OI: 10.7554/eLife.06416 Ku, et al., (2004). Conserved translational frameshift in dsDNA bacteriophage tail assembly genes. Molecular Cell 16, 11-2

Mousing Around with Bacteriophages: Using Antibodies Raised in BALB/c Mice to Analyze Phages

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