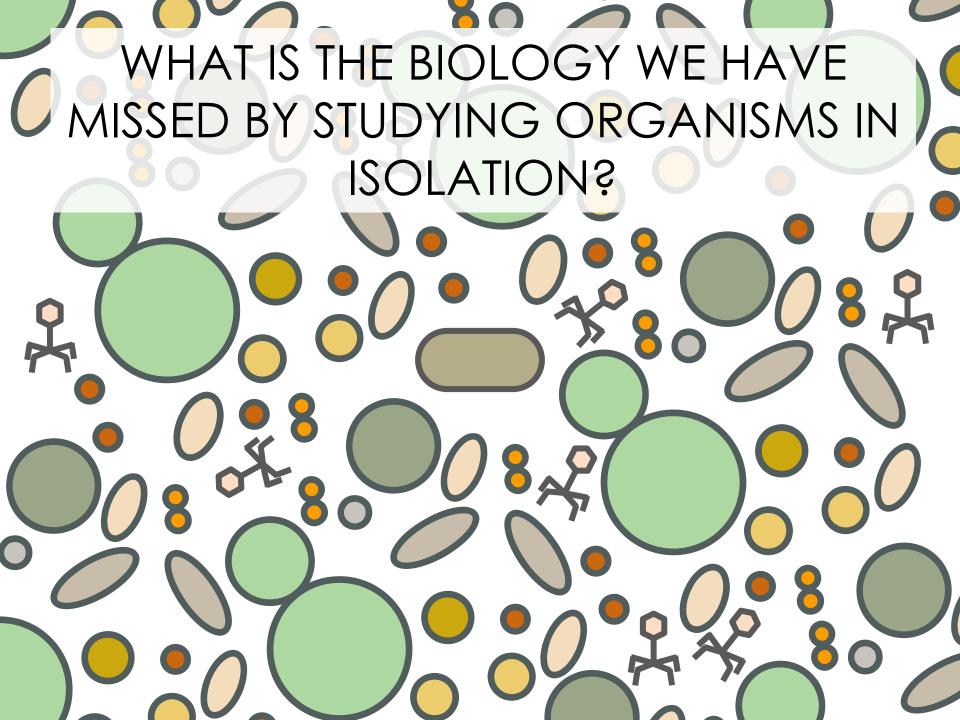
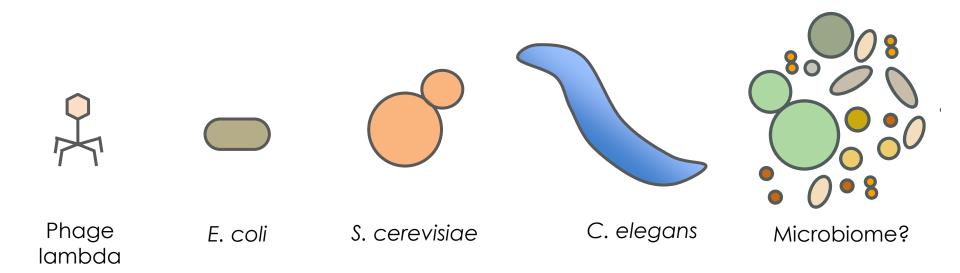
DISSECTING SPECIES INTERACTIONS IN THE CHEESE MICROBIOME

Rachel Dutton, PhD Associate Professor Division of Biological Sciences UC San Diego

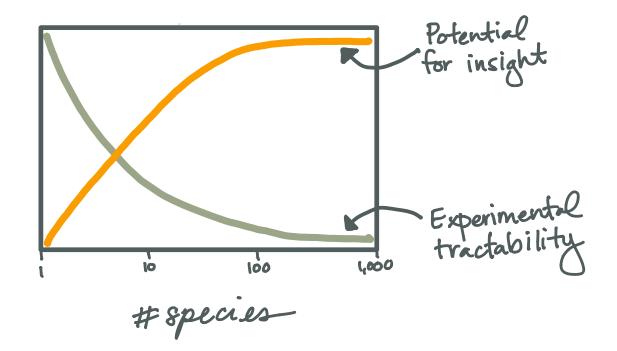


MODEL SYSTEMS ARE POWERFUL

Make observations \rightarrow Test hypotheses



BUT WHICH COMMUNITY?



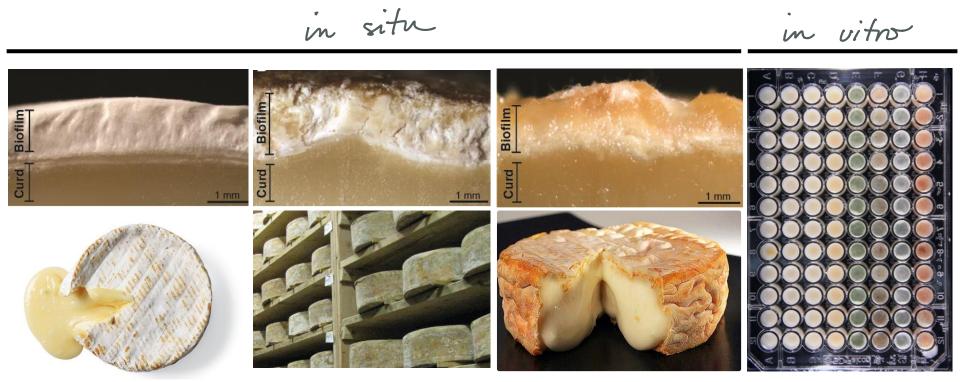
FERMENTED FOODS FALL IN THE "GOLDILOCKS ZONE"



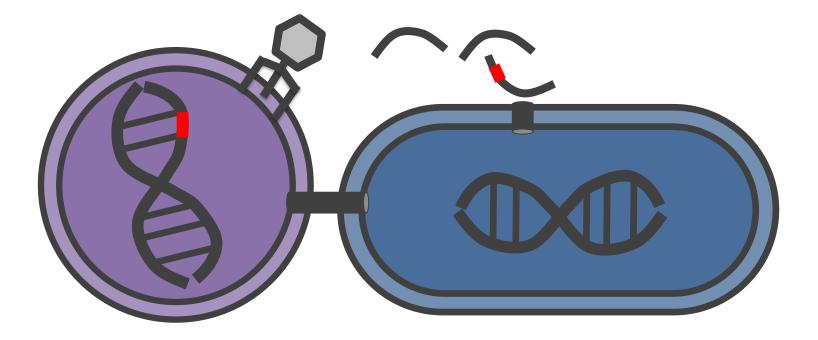
Wolfe & Dutton, Cell, 2015

THE CHEESE RIND BIOFILM

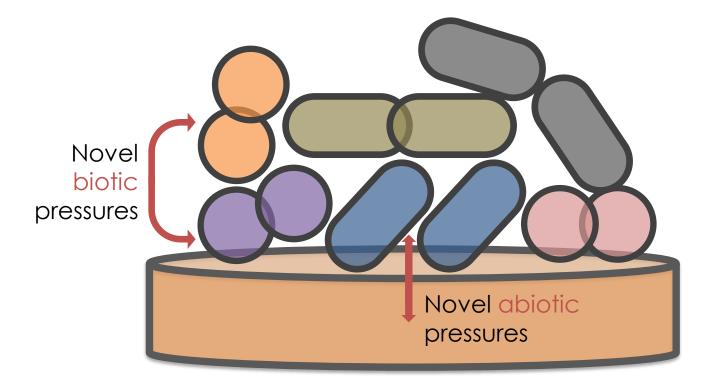
Communities range from low to medium complexity, phylogenetically diverse, completely culturable, amenable to *in vitro* reconstruction



HORIZONTAL GENE TRANSFER OCCURS WITHIN COMMUNITIES

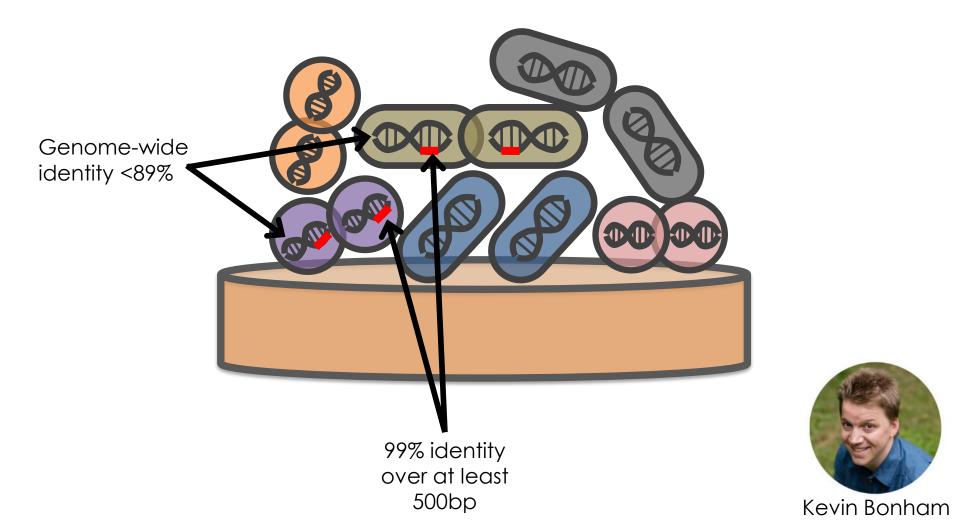




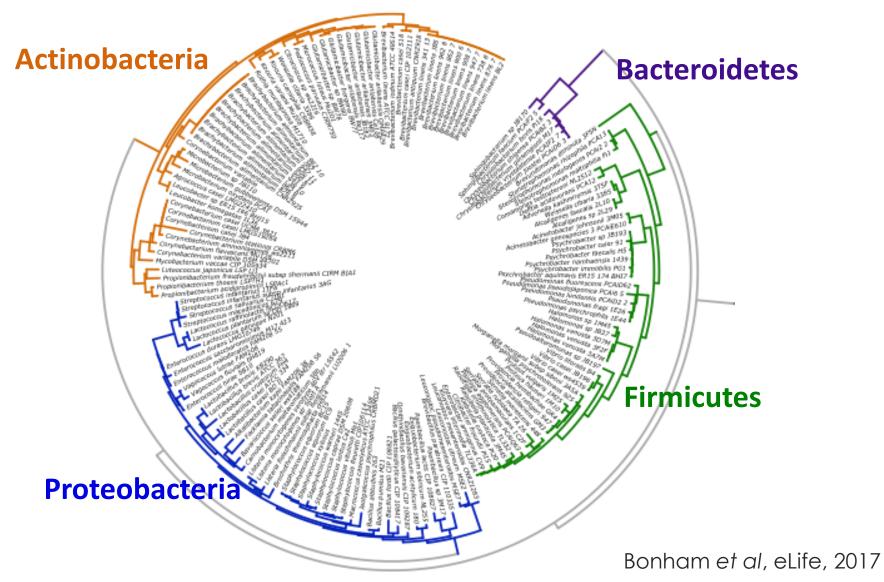


WHAT GENES ARE HORIZONTALLY TRANSFERRED?

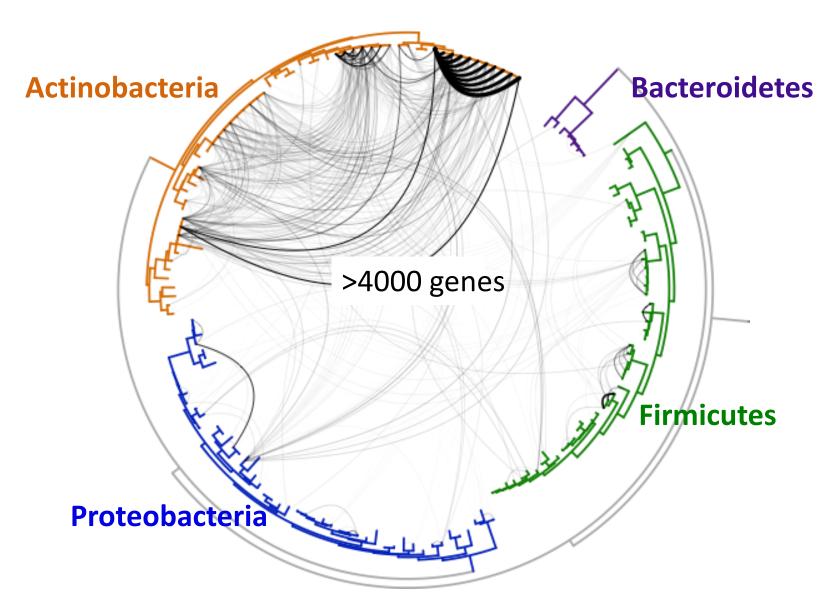
Looked for regions of high identity between pairs of genomes (Smillie et al, 2011)



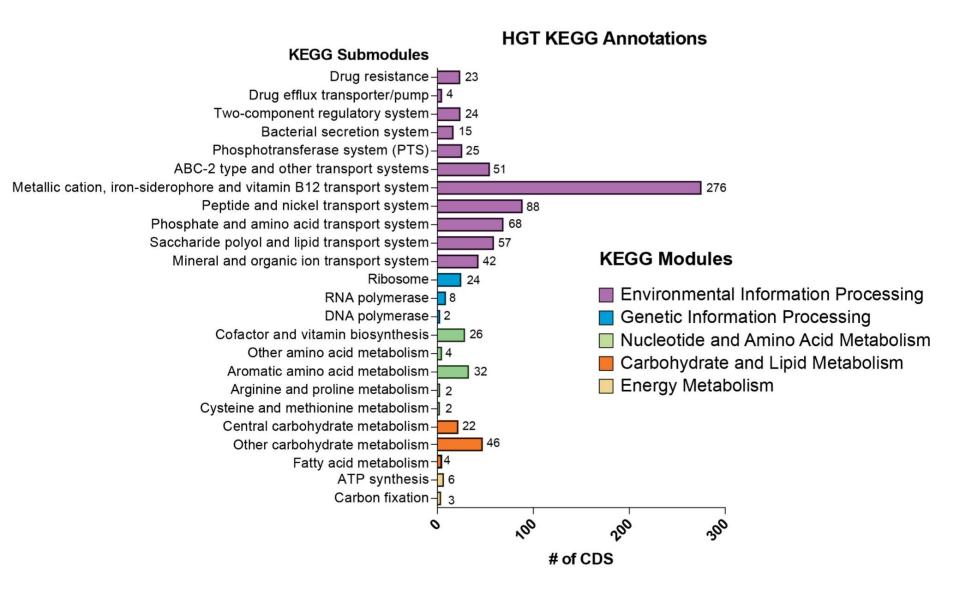
165 CHEESE-ASSOCIATED BACTERIAL GENOMES



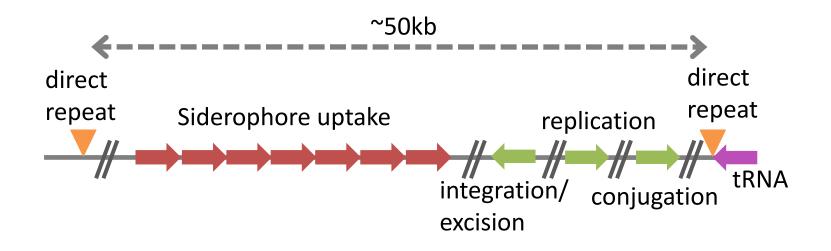
EXTENSIVE HGT IN CHEESE



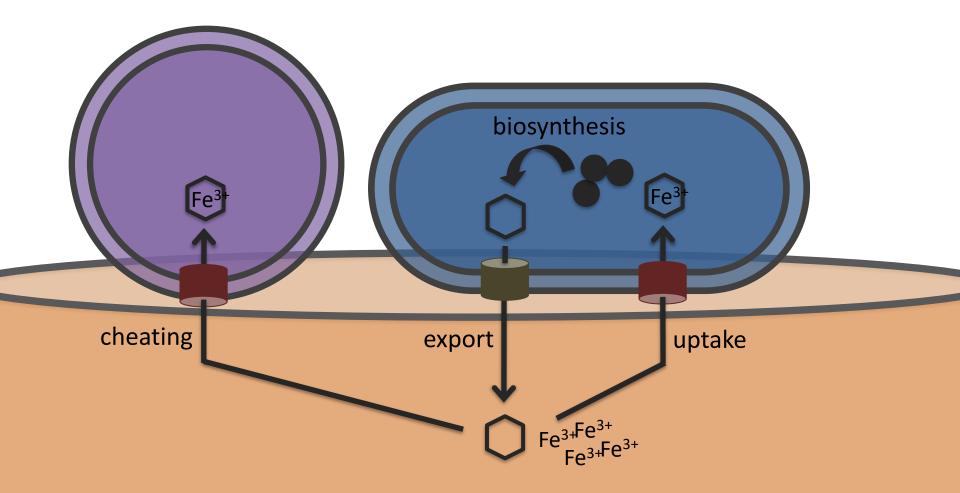
TRANSPORT SYSTEMS, ESPECIALLY FOR IRON, FOUND IN HGT REGIONS



SIDEROPHORE UPTAKE GENE CLUSTERS REPEATEDLY FOUND ON MOBILE ELEMENTS



POTENTIAL CHEATING IN CHEESE MICROBIOMES



WHAT IS THE DIVERSITY OF PLASMIDS AND PHAGE IN CHEESE RINDS (AND KEFIR)?

Sequencing: minION 9.4 Basecalling: guppy 4.0.15 Assembly: metaFlye Polishing: racon, medaka, DIAMOND/MEGAN plasmids: plasflow, PlasmidVerify, plasmidDB viruses: DeepVirFinder, VirSorter, ViPTree, vcontact2 Mobilization: mob-suite Visualization: anvi'o, bandange

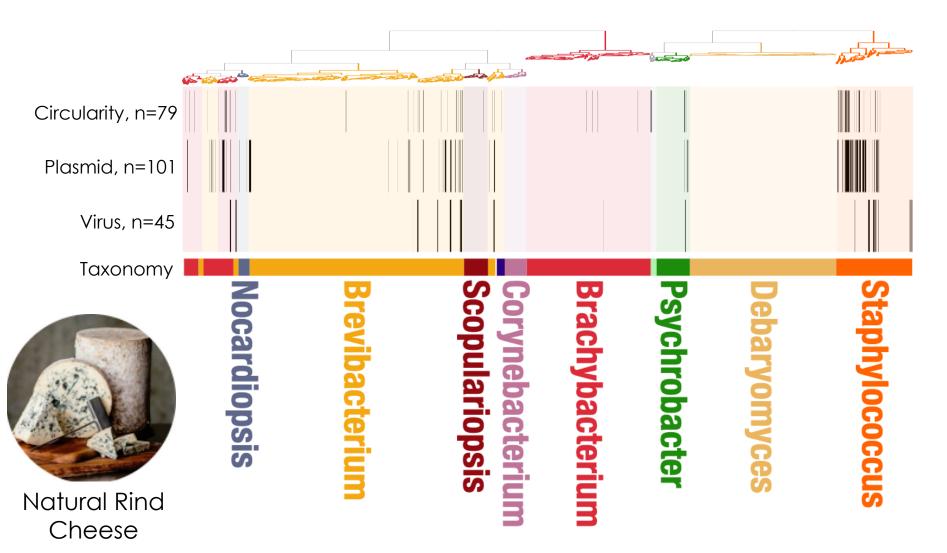
N ntact2

Cong Dinh

MD/PhD student

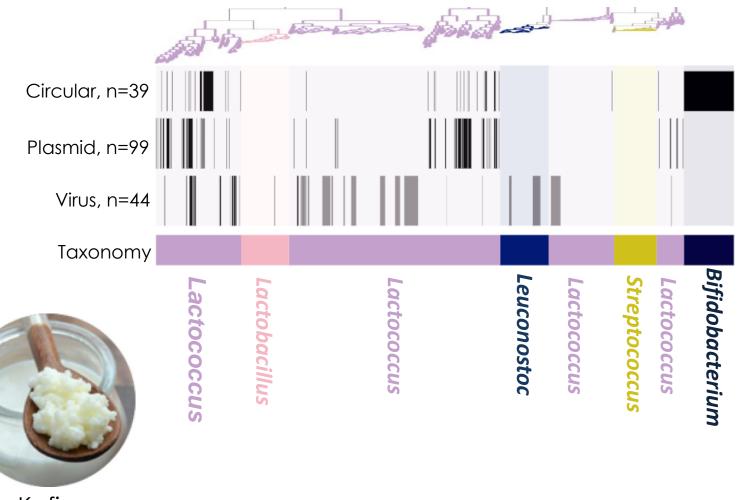
LONG-READ METAGENOMICS REVEALS ABUNDANT PLASMIDS AND VIRUSES

Clustering of contigs based on tetramer frequency and contig coverage (Anvi'o)



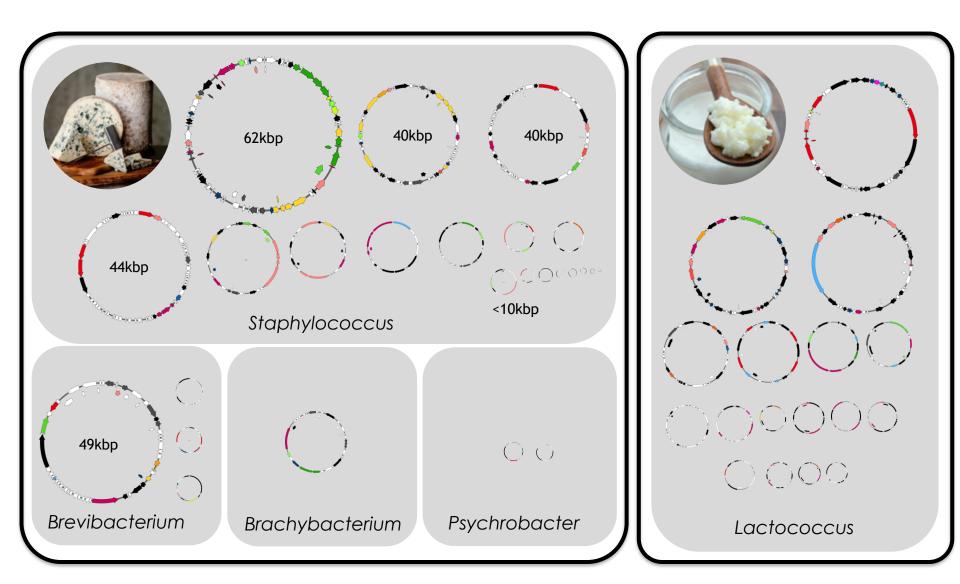
LONG-READ METAGENOMICS REVEALS ABUNDANT PLASMIDS AND VIRUSES

Clustering of contigs based on tetramer frequency and contig coverage (Anvi'o)



Kefir

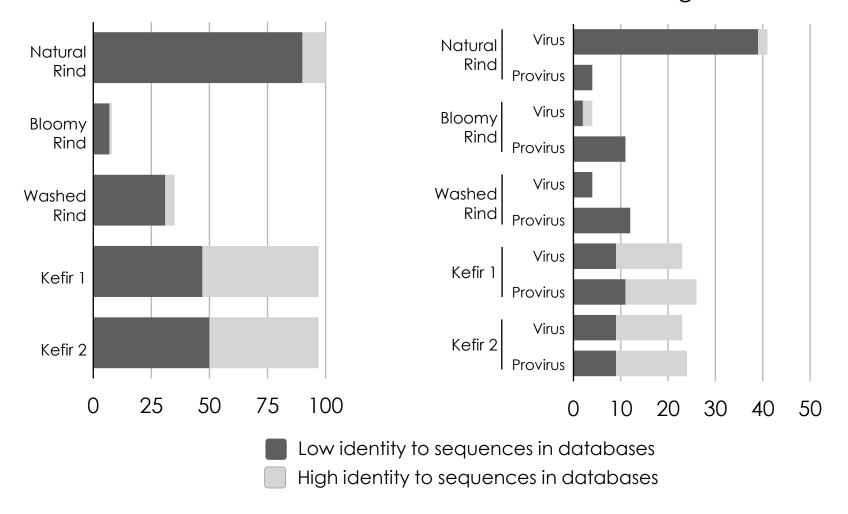
LONG READS ALLOWS ASSEMBLY OF CIRCULAR PLASMIDS



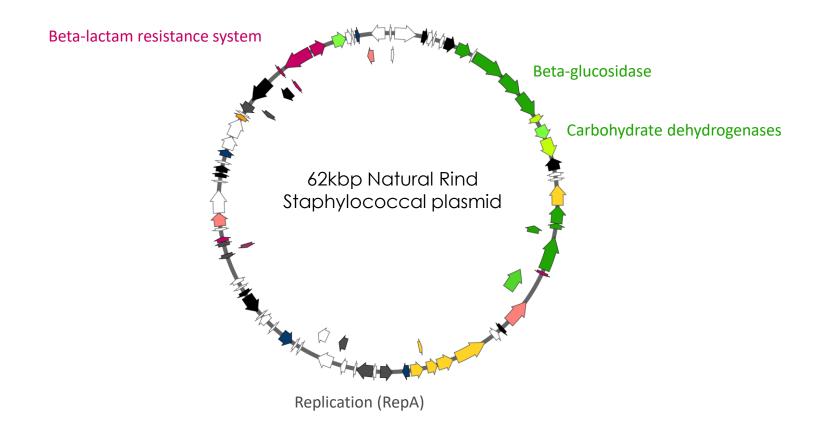
MANY NOVEL PLASMIDS AND PHAGE IN CHEESE RINDS

Plasmids

Phage



PLASMIDS MAY EXPAND HOSTS' CARBON ACCESS

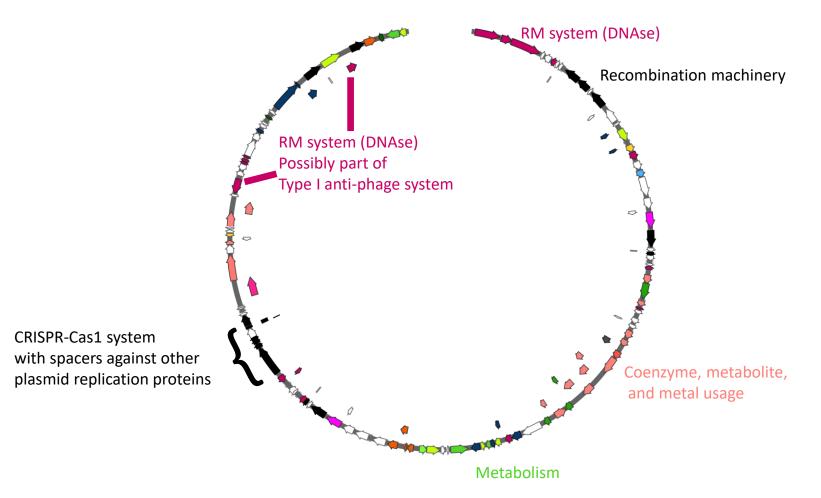


METAL RESISTANCE AND UPTAKE IS COMMON

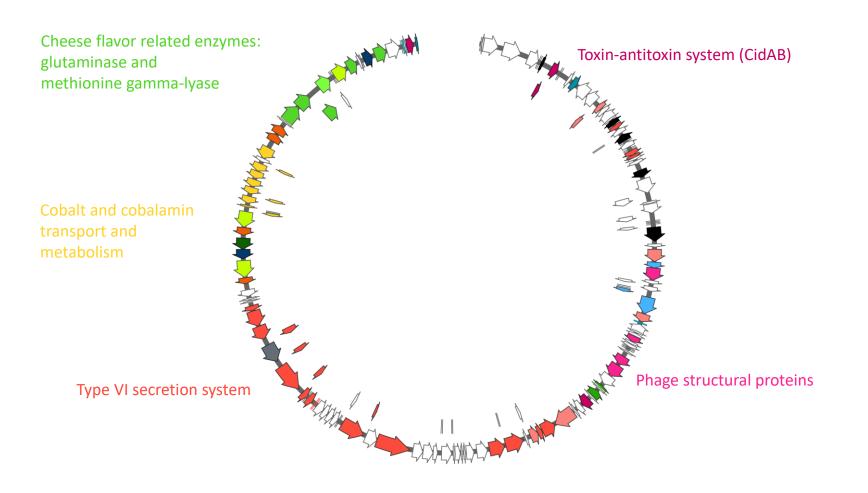
especially cadmium-resistance



DEFENSE PLASMID RESISTS OTHER PLASMIDS AND PHAGE



HAFNIA PROPHAGE ENCODING CHEESE FLAVOR GENES?





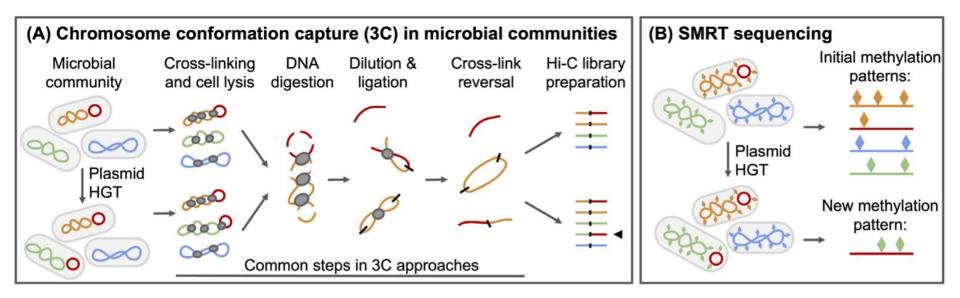
FEMS Microbiology Reviews, fuaa025, 44, 2020, 606–630

doi: 10.1093/femsre/fuaa025 Advance Access Publication Date: 16 July 2020 Review Article

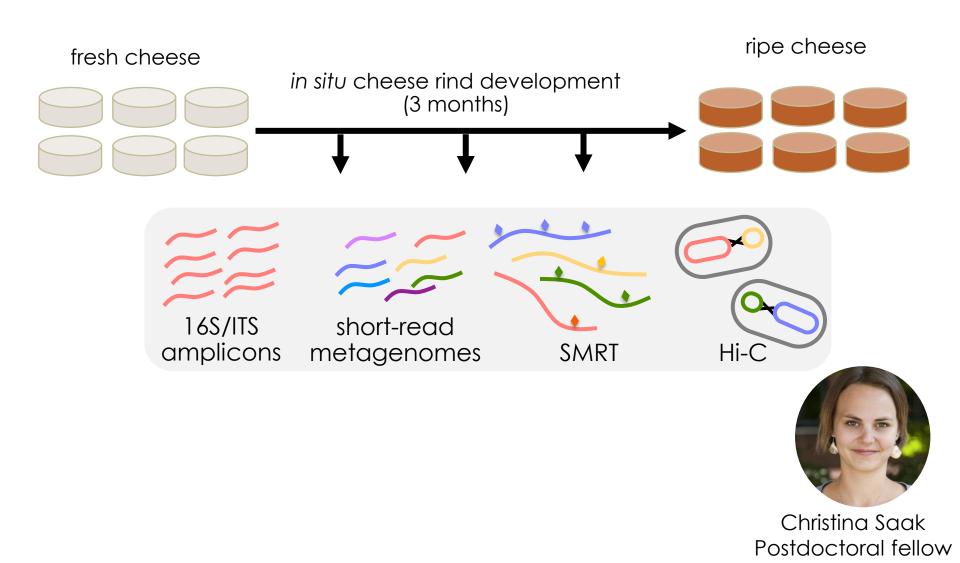
REVIEW ARTICLE

Experimental approaches to tracking mobile genetic elements in microbial communities

Christina C. Saak, Cong B. Dinh and Rachel J. Dutton*

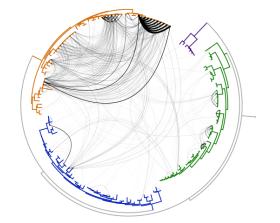


CAN WE OBSERVE HGT IN CHEESE?



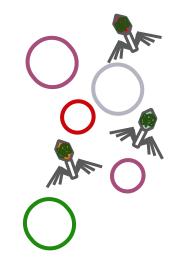
ABUNDANT EVIDENCE OF HGT IN CHEESE GENOMES

 REPEATED TRANSFER OF IRON UPTAKE PATHWAYS

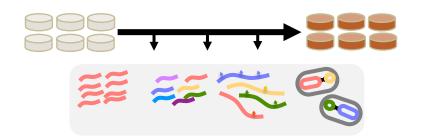


DIVERSE SET OF PLASMIDS AND PHAGE EVEN IN "SIMPLE" COMMUNITIES

SO MUCH MORE BIOLOGY TO
DISCOVER AND EXPLORE!



DEVELOPING APPROACHES TO TRACK HGT





CURRENT LAB MEMBERS

<u>Postdocs</u>: Manon Morin, Gary Heussler, Christina Saak, Emily Pierce <u>Graduate students</u>: Brooke Anderson, Cong Dinh, Tara Spencer <u>Undergraduates</u>: Brooke Johnson, Angel Sarabia, Martha Herrera <u>Lab manager</u>: Steven Villareal

FORMER LAB MEMBERS

<u>Postdocs</u>: Benjamin Wolfe (Tufts University), Julie Button (Seres Health), Kevin Bonham (Wellesley College) <u>Undergraduates</u>: Carla Espinoza, Val Jackson-Hundley, Gillian Belk, Antony Yang, Will Bushnell, Tyler Nelson, Kien Malarney, Daniel Rubin, Shanice Webster, Alyson Yee, Rajashree Mishra, Miriam Schiffman, Juan Alvarez, Adriann Negreros

COLLABORATORS

Laura Sanchez (UIC) Benjamin Wolfe (Tufts) Nick Loman (U Birmingham) Kit Pogliano (UCSD) Pieter Dorrestein (UCSD) Rob Knight (UCSD) Nancy Keller (UW-Madison) Sergey Kryazhimskiy (UCSD) Michael Harms (Univ of Oregon)

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