2020 Boston Bacterial Meeting

Day 1 - Thursday, July 16

9:00 AM	Opening Remarks		
I: Bacterial Physiology Session chair: Mona Orr			
9:10 AM	Felix Wong	Cytoplasmic condensation induced by membrane damage contributes to antibiotic lethality	
9:30 AM	Kanika Khanna	Asymmetric localization of the cell division machinery during Bacillus subtilis sporulation	
9:50 AM	Simone Pelliciari	Evidence of a broadly conserved bacterial chromosome origin unwinding system	
10:10 AM	Grace Johnson	Functionally Uncoupled Transcription-translation in B. subtilis	
10:30 AM		Break	
II: Antibiotic Resistance Session chair: Wendy Mo			
11:00 AM	Trever Smith	Morphological profiling to identify drug pathways of action in Mycobacterium tuberculosis	
11:20 AM	Melanie Martinsen	Core antibiotic-induced transcriptional signatures reflect susceptibility to all members of an antibiotic class	
11:40 AM	Anna Green	Genome-wide detection of epistasis in antibiotic resistant M. tuberculosis	
12:00 PM	Wenwen Huo	Novel molecular mechanisms for the development of fluoroquinolone resistance in vivo	
12:20 PM		Lunch	
1:30 PM		Breakout Sessions (Please see abstract book for links)	
2:30 PM		Break	
III: Phage a	Session chair: Batbileg Bor		
3:00 PM	Alex Meeske	A phage-encoded anti-CRISPR enables complete evasion of type VI-A CRISPR-Cas immunity	
3:20 PM	Kevin Gozzi	Caulobacter crescentus produces a gene transfer agent to mediate genetic exchange during stress	
3:40 PM	Siân Owen	Prophages have cognate immunity to their own phage resistance proteins	

Day 2 - Friday, July 17

Host Interactions	Session chair: Emrah Altindis
Alyson Warr	Shiga toxin shapes the intestinal innate immune response to EHEC
Mindy Engevik	Mucin degrading microbes release monosaccharides that chemoattract <i>Clostridioides</i> difficile and facilitate colonization of the intestinal mucus layer
Sage Foley	Microbiome regulation of intestinal P-glycoprotein expression in maintenance of homeostasis
Qian Huang	A gut commensal, Parabacteroides distasonis, triggers Type 1 diabetes autoimmunity
	Break
obial Interactions	Session chair: Joseph Larkin
Giuseppina Mariano	A new family of Type VI secretion system-delivered effector proteins displays ion-selective pore-forming activity
Rebecca McHugh	Characterisation of aurodox: A Type III secretion system inhibitor from <i>Streptomyces</i> goldiniensis
Betty Slinger	Insights into the role of inter-species quorum sensing during co-infections with <i>B. multivorans</i> and <i>P. aeruginosa</i>
Vijay Singh	Anti-quorum sensing and anti-biofilm activity of 1,2-benzenedicarboxylic acid, diisooctyl ester against <i>Pseudomonas aeruginosa</i> by diminishing the quorum sensing-controlled virulence expression
	Lunch
hods in Microbiology	Session chair: Benjamin Ross
Bimal Jana	CRISPRi-TnSeq: A genome-wide high-throughput tool for bacterial essential and nonessential genetic interaction mapping
Jason Lynch	Development of designer probiotics for targeted delivery of immunomodulatory payloads
Sydney Blattman	Prokaryotic single-cell RNA sequencing by in situ combinatorial indexing
	Keynote Address - KC Huang
	Closing Remarks
	Mindy Engevik Sage Foley Qian Huang Tobial Interactions Giuseppina Mariano Rebecca McHugh Betty Slinger Vijay Singh Thods in Microbiology Bimal Jana Jason Lynch