

Welcome! The NIDA Genetics and Epigenetics Cross-Cutting Research Team (GECCRT) is a multi-institute working group with members from the intramural and extramural program staff from NIDA, NIAAA and NIMH. The goal of the GECCRT is to support the use of genetic and epigenetic tools to uncover new genetic, biochemical, and epigenetic pathways that contribute to substance use disorders. There is no registration. Webex links for each day are located in the agenda. The attendee password is: GECCRT21.

Day 1	March 8, 2021 Password: GECCRT21
Link:	https://nih.webex.com/nih/onstage/g.php?MTID=ea4943eef2b925a75ceeacf7b36534310
10:45	Welcome! Dr. Nora Volkow, Director National Institute on Drug Abuse
11:00 to	Session 1: Social Determinants of Health as Mediators of Phenotypes Relevant to SUD (Udi
12:30	Ghitza and Kerry Ressler, Session Chairs)  Goals: Interactions among genetic/epigenetic factors and social determinants of health may
	in some cases be mediators of phenotypes relevant to substance use disorders (SUD) and
	concomitant mood disorders. Speakers will critically discuss recent research on interactions of life stress or related adverse consequences with genetic and epigenetic factors mediating vulnerability for, or resilience to, substance use disorders and co-occurring anxiety-related
	conditions.
11:00	Kerry Ressler, PhD
	McLean Hospital, Harvard Medical School
	Introductory talk providing background for later presentations and research on interactions between social stress and genetic factors mediating vulnerability for post-traumatic stress disorder
11:15	
11:13	Christopher Guevara, Graduate Student Icahn School of Medicine at Mount Sinai
	Genetic mutations which may contribute to synaptic plasticity in brain areas affecting resilience to social stress
11:30	Jean Lud Cadet, MD
	NIDA Intramural Research Program
	Epigenetic and transcriptional consequences of compulsive methamphetamine taking and abstinence in the presence of adverse consequences
11:45	Shareefa Dalvie, PhD
	University of Cape Town, South Africa
	Genomic influences on self-reported childhood maltreatment and their relation to co- occurrence of psychiatric disorders
12:00	Discussion
12:30	Break

1:00	Session 2: Emerging Methods to Identify Missing Heritability
1:00	Randall J. Ellis, Graduate Student Icahn School of Medicine at Mount Sinai Genomic prediction of alcohol and opioid use disorders using machine learning
1:20	Seon-Kyeong Jang, Graduate Student University of Minnesota Rare variant heritability of tobacco use: Evidence from deep whole-genome sequencing of up to 26,000 individuals
1:40	Daniel Jacobson, PhD Oak Ridge National Laboratory Embracing Complexity: Supercomputing Enabled Systems Biology to Understand Complex Neurosystems
2:00	Slack Poster Sessions (drugabuse.slack.com)
5:00	Adjourn

Day 2	March 9, 2021 Password: GECCRT21
Link:	https://nih.webex.com/nih/onstage/g.php?MTID=ef20bbcc69ac66938d988d067ff032d89
10:45 AM	Welcome and Summary
11:00 to	Session 3: Novel Approaches for Target Discovery and Therapy for Substance Use Disorders
12:30	(Jonathan Pollock, Session Chair)
	This session highlights genetic approaches to identify novel target and therapeutics for
	substance use disorders.
11:00	Rohan Palmer, PhD
	Emory University Using Inverse Transcriptomics to identify treatments for SUD
11.15	
11:15	Brock Grill, PhD The Scripps Research Institute
	Genetic behavioral screen identifies an orphan anti-opioid system
11:30	Vivek Kumar, PhD
	The Jackson Laboratories
	Identifying potential drug targets from the KOMP mice
11:45	Michelle Hastings, PhD
	Rosalind Franklin Medical University
	Antisense approaches to the treatment of Disease
12:00	Discussion
12:30	Break
1:00	Session 4: The Microbiome in Substance Use Disorders
1:00	Christiann Gaines, Graduate Student
	University of North Carolina, Chapel Hill
	Assessing the impact of compositionally distinct gut microbiotas on differences in initial
	cocaine sensitivity in closely related inbred mouse substrains
1:20	Sierra Simpson, PhD
	University of California, San Diego Drugs, Bugs, and SUDs: Leveraging the Microbiome and Metabolome to Predict Addiction
	Liability
1:40	Drew Kiraly, MD, PhD

2:00 Slack Poster Sessions (drugabuse.slack.com) 5:00 Adjourn  Day 3 March 10, 2021 Password: GECCRT21 Link: https://nlin.webex.com/nih/onstage/g.php?MTID=ece26a4f78866bb6178c91673f0869ecb  10:45 Welcome and Summary  11:00 to Session 5: Beyond GWAS: Mobile DNA in Disease (Amy Lossie, Session Chair)  The purpose of this session is to introduce the SUD Genomics field to strategies that assess the parts of the genome that are typically not included in GWAS studies: transposable elements and structural variations.  11:00 Alex Urban, PhD Stanford University Analysis of Line1 elements and Structural Variants in the brain  11:15 Melissa Gymrek, PhD University of California, San Diego The Impact of Short Tandem Repeats on Gene Expression  11:30 Avi Nath, MD National Institute of Neurological Disorders and Stroke Intramural Research Program Regulation of stem cell function and neuronal differentiation by HERV-K via mTOR pathway  11:45 Sandhya Chandrasekaran, MSTP Student Icahn School of Medicine at Mount Sinai Cell-type specific chromatin configuration in the mammalian brain highlights a novel organizational architecture of an active class of mobile elements, the endogenous retroviruses, in neurons  12:00 Discussion  12:30 Break  1:00 Session 6: The Interplay Between Substances of Abuse and Gene Regulation  Jimmy Olusakin, PhD University of Maryland School of Medicine Transcriptomic adaptations in emotional and sensory brain nuclei in perinatal fentanyl exposed rodents  4:20 Marta Pratelli, PhD University of California, San Diego Drugs of abuse drive activity producing changes in gene expression that switch neurotransmitters and behaviors  4:40 Jian Feng, PhD Florida State University Neuron-specific role of methylated DNA cytosine dioxygenase TET1 in cocaine addiction  5:00 Wrap Up and Adjourn		Icahn School of Medicine at Mount Sinai  Manipulations of gut microbiome diversity alter cocaine seeking behavior and striatal gene expression
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