



## AGENDA

### ***Day 0 – Wednesday, 8/21***

<i>Time</i>	<b>Event and Description</b>	<b>Location</b>
<b><i>7:30 – 10:30 PM</i></b>	Networking Session - Night out (informal)	Tommy Joe's

### ***Day 1 – Thursday, 8/22***

<i>Time</i>	<b>Event and Description</b>	<b>Location</b>
<b><i>7:30 – 8:30 AM</i></b>	Registration opens, coffee provided	Welcome Area
<b><i>8:30 – 8:45 AM</i></b>	Welcome session – Introductions (JI Committee & NCI)	Auditorium
<b><i>8:45 – 9:15 AM</i></b>	Patient Advocates Session: Hearing their story	Auditorium
<b><i>9:15 – 10:15 AM</i></b>	Keynote: Dr. Hadi Nia	Auditorium
<b><i>10:15 – 10:30 AM</i></b>	Mental Health Break	
<b><i>10:30 – 12:00 AM</i></b>	Session I: Podium Talks with Patient Advocates - Tumor immunology and microenvironment <ul style="list-style-type: none"> <li>• Pablo Llevenen: Plasma exosomes in obesity-driven diabetes exacerbate progression of triple negative breast cancer: Insights from animal models</li> <li>• Kun Han: Spatial analysis of immune-related adverse events using novel animal models in myocarditis and neurotoxicity</li> <li>• Cassidy Hagan: Apoptotic cells promote circulating tumor cell survival and metastasis</li> <li>• Junior West: Claudin 7 suppresses invasion and metastasis through repression of the smooth muscle actin cytoskeleton</li> <li>• Joshua Ginzel: Dynamic modeling of tumor progression illuminates an occult transition that establishes metastatic potential</li> </ul>	Auditorium

<b>12:00 – 1:30 PM</b>	Lunch Special Sessions: <ul style="list-style-type: none"> <li>• Scalable Processing of Multiplexed Tissue Imaging Datasets Using Nextflow and MCMICRO (pre-registration required)</li> <li>• NCI SBIR: Small Business Transition Grant Info Session</li> </ul>	Welcome Area  Rooms F1/F2  Rooms E1/E2
<b>1:30 – 2:30 PM</b>	Concurrent Session IIA: Spotlight Talks - Translational cancer research and drug discovery <ul style="list-style-type: none"> <li>• Samson Eugin Simon: DNA hypermethylation impacts tumor suppressor genes in BXD preclinical mouse models</li> <li>• Kacey Ronaldson-Bouchard: Recapitulation of skeletal metastasis using an in vitro organ- on-a-chip platform</li> <li>• Zhengyuan Pang: In situ visualization of covalent cancer drug binding in 3D mammalian tissue</li> <li>• Wenjuan Dong: Enhancing prediction of trastuzumab deruxtecan response and resistance mechanisms in metastatic breast cancer through spatial profiling and cellular crosstalk analysis</li> <li>• Lucas ZhongMing Hu: Elucidating oncology drug mechanism of action and polypharmacology with a large-scale perturbational profile compendium</li> <li>• Yuhan Qiu: Insulin resistance increases TNBC aggressiveness and brain metastasis via adipocyte-derived exosomes</li> </ul>	Balcony B
<b>1:30 – 2:30 PM</b>	Concurrent Session IIB: Spotlight Talks - Multiomics and bioinformatics in cancer research <ul style="list-style-type: none"> <li>• Behnaz Bozorgui: Single cell spatial proteomic analysis and computational evaluation pipeline</li> <li>• Kyle MacQuarrie: Chromosome specific organizational characteristics in rhabdomyosarcoma and normal myogenic cells</li> <li>• Pascal Belleau: Computational framework for inference of genetic ancestry from challenging human molecular data</li> <li>• Reshma Kalyan Sundaram: Integrating signaling and transcription to study c-Myc induced stress response in cancerous cells</li> <li>• Seongyeol Park: Organ-specific evolution of tumor microenvironments in metastatic</li> </ul>	Balcony C

	<p>breast cancer</p> <ul style="list-style-type: none"> <li>• Kaiyuan Zhu: Elucidating genomic and three-dimensional architectures of extrachromosomal DNA</li> </ul>	
<b>2:30 – 3:30 PM</b>	Networking Session: Speed Dating	Atrium
<b>3:30 – 4:30 PM</b>	<p>Concurrent Session IIIA: Spotlight Talks - Niche regulation of cancer progression</p> <ul style="list-style-type: none"> <li>• Christina Ennis: Plasma exosomes from individuals with type 2 diabetes drive breast cancer progression in patient-derived organoids</li> <li>• Ethan Seltzer: Neuronal substance-P drives breast cancer growth and metastasis</li> <li>• Lucy Britto: Lymphoma organoids reveal T cells spatially alter B cell receptor signaling via histone modifications</li> <li>• Ghmkin Hassan: Antidepressants inhibit pancreatic ductal adenocarcinoma progression by reducing fibrosis and perineural invasion</li> <li>• Anna Kolarzyk: Investigating VEGF-A induced lymphatic vessel remodeling and its implications for immune cell trafficking in pancreatic cancer</li> <li>• Toshiro Hara: In situ state of invading glioblastoma cells by single-cell and spatial transcriptomics</li> </ul>	Balcony B
<b>3:30 – 4:30 PM</b>	<p>Concurrent Session IIIB: Spotlight Talks - Genetic and epigenetic regulation of tumor progression and metastasis</p> <ul style="list-style-type: none"> <li>• Nana Adjoa Ben-Crentsil: RNA interference of P65 activity mediates oncogenic inflammation in TET2 mutated clonal hematopoiesis</li> <li>• Sarah Groves: Modeling Phase Separation of a Mitosis Signaling Network in Breast Cancer</li> <li>• Carolina De Santiago: Analyzing the clonal dynamics and metastatic potential of isolated subpopulations within TNBC cells</li> <li>• Monika Dhankhar: Dynamics of chromatin reorganization in response to chemo-mechanical environmental cues</li> <li>• Lina Kroehling: High-resolution characterization of age-specific changes in the HPV-negative HNSCC tumors</li> <li>• Tatiana Miti: Combining spatial analysis</li> </ul>	Balcony C

	methods for studies of stromal effects on tumor remission-relapse dynamics	
<b>4:30 – 5:30 PM</b>	Poster Session I	Atrium
<b>6:00 – 7:00 PM</b>	Depart for The Bethesda Hotel (shuttle provided)	Shuttle
<b>7:00 – 9:00 PM</b>	Dinner	The Bethesda Hotel
<b>9:00 – 11:00 PM</b>	Networking Session: Sightseeing (informal)	Bethesda and Washington, D.C.

## **Day 2 – Friday, 8/23**

<b>Time</b>	<b>Event and Description</b>	<b>Location</b>
<b>8:00 – 9:00 AM</b>	Registration opens, coffee provided	Welcome Area
<b>9:00 – 10:30 AM</b>	Special Session I: What can you do with a PhD? Breakfast served	Rooms A/B and E1/E2
<b>10:30 – 12:00 AM</b>	Session IV: Podium Talks with Patient Advocates - Multiomics and Computational Cancer Research <ul style="list-style-type: none"> <li>• Dig Vijay Kumar Yarlagadda: Discrete representation learning for modeling imaging-based spatial transcriptomics data</li> <li>• Minjeong Kim: Unveiling genetic complexity of lung adenocarcinoma by identifying a novel EGFR mutation signature</li> <li>• Mohd Saqib: Investigating tumor immune suppression through integration of PD1-PD-L1 interactions with MDSC &amp; T cell phenotypes</li> <li>• Riley Manning: 3D agent-based modeling of glioblastoma subtypes and therapies</li> <li>• Vikas Pandey: AI-enhanced rapid lifetime determination method for fast macroscopic fluorescence lifetime imaging</li> </ul>	Auditorium
<b>12:00 – 1:30 PM</b>	Lunch Special Sessions: <ul style="list-style-type: none"> <li>• NCI CBIIT Data Science Training Resources</li> </ul> Programming Booths: SBIR, BTEP, & CCT	Welcome Area  Rooms F1/F2; Overflow G1/G2
<b>1:30 – 3:00 PM</b>	Special Session II: Just Starting Out Panel	Auditorium
<b>3:00 – 4:30 PM</b>	Poster Session II	Atrium
<b>4:30 – 5:30 PM</b>	Concurrent Session VA: Spotlight Talks - AI/ML approaches for translational cancer research <ul style="list-style-type: none"> <li>• Sehyun Oh: OmicsMLRepo: Ontology-leveraged metadata harmonization to improve AI/ML-readiness of omics data in Bioconductor</li> </ul>	Balcony B

	<ul style="list-style-type: none"> <li>• Yi Lian: A reliable and adaptive framework for high-dimensional fairness aware integration of multiple datasets</li> <li>• Hairong Wang: A self-supervised AI framework for quantitative assessment of intra-tumoral heterogeneity in GBM using MRI</li> <li>• Sushant Patkar: Predicting the tumor microenvironment molecular composition from histopathology images to characterize immunotherapy responses in non-small cell lung cancer patients</li> <li>• Xinling Li: Mathematical model of AND-gated protease sensors predicts tunable improvements to signal-to-noise</li> <li>• Alexander Lachman: ARCHS4 navigator: multimodal feature vectors as descriptors of gene expression data</li> </ul>	
<b>4:30 – 5:30 PM</b>	<p>Concurrent Session VB: Spotlight Talks - Tumor immunology and immunotherapy</p> <ul style="list-style-type: none"> <li>• Ali Zamat: Image-guided sonogenetic control of thermal-sensitive CAR T cells overcomes brain tumor heterogeneity and immune suppression</li> <li>• Kolade Adebawale: Dynamics of macrophage tumor infiltration</li> <li>• Larry Dooling: Macrophage clusters extend multiple pseudopods to disrupt cell junctions between cohesive targets</li> <li>• Michelle Loui: Autoantibodies in high-grade serous ovarian cancer interact poorly with cytotoxicity-inducing Fc receptors</li> <li>• Udochi Felicia Azubuike: Microenvironmental regulation of immune cell infiltrates in cutaneous melanoma</li> <li>• Yufei Cui: Quantitative cell type specific immunopeptidome analysis during macrophage and tumor co-evolution reveals therapeutic MHC-I restricted peptides in glioblastoma</li> </ul>	Balcony C
<b>5:30 – 6:00 PM</b>	Closing Session	Auditorium
<b>6:00 PM</b>	Conference Adjourns	