

**Cancer Steering Committee  
 Annual Scientific Symposium**  
*Bethesda Marriott Hotel*  
*Bethesda, MD*

Monday, November 4, 2019 11:00 am – 8:00 pm  
 Tuesday, November 5, 2019 7:30 am – 4:00 pm

**Agenda**

<b>Monday, November 4</b>	
<b>11:00 am – 12:00 pm</b>	<b>Registration and Lunch</b>
<b>12:00 – 12:15 pm</b>	<b>FNIH Welcome</b> – Stacey Adam and Dana Connors
<b>12:15 – 12:45 pm</b>	<b>Meeting Introduction and Overview</b> – Eric Rubin, Merck and Gary Kelloff, NCI
<b>12:45 – 1:15 pm</b>	<b>Keynote</b> – Monica Bertagnolli, Dana Farber Cancer Institute
<b>1:15 – 2:45 pm</b>	<p><b>Biomarkers in the Immunotherapy Landscape</b>          Chair: Eric Rubin, Merck</p> <ul style="list-style-type: none"> <li>• <b>CRI Landscape Analysis of IO Trials and Biomarkers</b>              Jun Tang, Cancer Research Institute</li> <li>• <b>Human Tumor Atlas</b>              Bruce Johnson, Dana Farber Cancer Institute</li> <li>• <b>Perspectives on Biomarkers in IO Trials</b> <ul style="list-style-type: none"> <li>– Priti Hegde, Foundation Medicine</li> <li>– Lyndsay Harris, NCI/Partnership for Accelerating Cancer Therapies</li> <li>– Theresa LaVallee, Parker Institute for Cancer Immunotherapy</li> <li>– Alessandra Cesano, Essa Pharma</li> </ul> </li> <li>• <b>Discussion/Panel</b></li> </ul>
<b>2:45 – 3:00 pm</b>	<b>Coffee/Networking</b>

3:00 – 4:30 pm	<p><b>Challenges for Biomarker Development: Heterogeneity, Immunotherapy-related Toxicity, and Regulatory Approaches</b> Chair: Bruce Johnson, Dana Farber Cancer Institute</p> <ul style="list-style-type: none"> <li>• <b>ctDNA in Immuno-oncology</b> Alex Snyder, Merck</li> <li>• <b>Data-Driven Evidence of Intra- and Inter-tumor Heterogeneity</b> Kurt Schalper, Yale</li> <li>• <b>Immunotherapy-related Adverse Events—Infrastructure for Evaluation</b> Elad Sharon, NCI</li> <li>• <b>Regulatory Perspective</b> Marc Theoret, FDA</li> <li>• <b>Discussion/Panel</b></li> </ul>
4:30 – 4:45 pm	Coffee/Networking
4:45 – 6:30 pm	<p><b>Advanced Technologies in Biomarker Development</b> Chair: Greg Friberg, Amgen</p> <ul style="list-style-type: none"> <li>• <b>Proteotyping Human Cancers</b> Dan Liebler, Protypia</li> <li>• <b>The SCANT Method</b> Jonathan Carlson, Harvard</li> <li>• <b>Diagnostic Technologies to Guide Therapy Design and Follow Therapy Response in Immuno-oncology</b> James Heath, Institute for Systems Biology</li> <li>• <b>Epigenetic Signatures in the Immune Tumor Interface: Stratifying Melanoma Patients for Checkpoint Inhibitors</b> Claudio Carini and Peter Parker, King's College London</li> <li>• <b>Discussion/Panel</b></li> </ul>
6:30 – 7:00 pm	Keynote – Chi Van Dang, University of Pennsylvania
7:00 – 8:00 pm	Reception
8:00 pm	Adjourn – Dinner on Your Own

## AGENDA: TUESDAY, NOVEMBER 5

<b>Tuesday, November 5</b>	
<b>7:30 – 7:50 am</b>	<b>Registration and Breakfast</b>
<b>7:50 – 8:00 am</b>	<b>Meeting Overview II – Eric Rubin, Merck</b>
<b>8:00 – 8:30 am</b>	<b>Keynote – Chris Boshoff, Pfizer</b>
<b>8:30 – 8:45 am</b>	<b>Break/Networking</b>
<b>8:45 – 10:15 am</b>	<p><b>Breakout Session A – The Microbiome</b> Chair: Yasmine Belkaid, NIAID and NIH Center for Human Immunology</p> <ul style="list-style-type: none"> <li>• <b>An Overview of the Science of the Microbiome</b> Yasmine Belkaid, NIAID</li> <li>• <b>Functional Microbiomic Signatures as Predictors of Pro-tumor Inflammatory State and Susceptibility to Immunotherapy</b> Leopoldo Segal, NYU</li> <li>• <b>Discussion/Panel</b></li> </ul>
<b>8:45 – 10:15 am</b>	<p><b>Breakout Session B – Measurable Residual Disease</b> Chair: Greg Reaman, FDA</p> <ul style="list-style-type: none"> <li>• <b>Statistical Considerations Regarding the Role of MRD</b> Don Berry, Berry Consulting</li> <li>• <b>Molecular Response and MRD in AML</b> Jerry Radich, University of Washington</li> <li>• <b>The Future of MRD in Multiple Myeloma</b> Ken Anderson, Dana Farber Cancer Institute</li> <li>• <b>Next Steps for MRD in Other Hematologic Cancers – AML</b> Robert Loberg, Amgen</li> <li>• <b>The Need for an Opportunity for MRD Standardization in AML</b> Kim Jessup, Unaffiliated</li> <li>• <b>Discussion/Panel</b></li> </ul>
<b>10:15 – 10:30 am</b>	<b>Coffee/Networking</b>

10:30 am – 12:25 pm	<p><b>Breakout Session C – Imaging Biomarkers</b> Chair: Annette Schmid, Takeda</p> <ul style="list-style-type: none"> <li>• <b>Imaging Biomarkers as Trial Endpoints</b> Greg Goldmacher, Merck</li> <li>• <b>Parameters Based on Tumor Growth and Regression</b> Tito Fojo, Columbia University</li> <li>• <b>Vol-PACT Project Read Out</b> Larry Schwartz, Columbia University</li> <li>• <b>Radiomic Feature Identification</b> Andrea Thiele and Bushi Wang, Boehringer Ingelheim</li> <li>• <b>Whole Organ Radiomics in Oncology: A Novel AI Based Approach</b> Nozha Boujemaa, Median Technologies</li> <li>• <b>Discussion/Panel</b></li> </ul>
10:30 am – 12:25 pm	<p><b>Breakout Session D – Liquid Biopsy</b> Chair: Carl Barrett, AstraZeneca</p> <ul style="list-style-type: none"> <li>• <b>Challenges in Establishing ctDNA Controls</b> Bob McCormack, Unaffiliated</li> <li>• <b>Standardization of Tumor Mutational Burden</b> Mark Stewart, Friends of Cancer Research</li> <li>• <b>Use of the Immunosequencing Platform with Cell-Free DNA</b> Lanny Kirsch, Adaptive Biotechnologies</li> <li>• <b>PD Assay Network and Other Collaborations</b> Howard Scher, Memorial Sloan Kettering Cancer Center</li> <li>• <b>Clinical Implementation Challenges</b> Stanley Hamilton, MD Anderson</li> <li>• <b>Collaborations and Initiatives to Validate Liquid Biopsy</b> <ul style="list-style-type: none"> <li>– Lauren Leiman, BloodPAC</li> <li>– Reena Philip, FDA</li> </ul> </li> <li>• <b>Discussion/Panel</b></li> </ul>
12:25 – 12:50 pm	<b>Lunch/Networking</b>
12:50 – 1:10 pm	<p><b>Lunch Session: Breakout Session Summaries</b></p> <ul style="list-style-type: none"> <li>• <b>The Microbiome</b> Session Chair: Yasmine Belkaid, NIAID</li> <li>• <b>Measurable Residual Disease</b> Session Chair: Greg Reaman, FDA</li> <li>• <b>Imaging Biomarkers</b> Session Chair: Annette Schmid, Takeda</li> <li>• <b>Liquid Biopsy</b> Session Chair: Carl Barrett, AstraZeneca</li> </ul>

1:10 – 1:30 pm	<b>Lunch Session: Pediatric Oncology Perspective</b> <ul style="list-style-type: none"> <li>• <b>Pediatric Oncology Patient Advocate</b> Caitlyn Barrett, CureSearch</li> <li>• <b>Clinical Trial Access and Inclusion/Exclusion Criteria</b> Ann Ramer, Advocate</li> </ul>
1:30 – 2:00 pm	<b>Keynote – Gideon Blumenthal, FDA</b>
2:00 – 3:50 pm	<b>Data Science and Machine Learning in Biomarker Development</b> Chair: Hisham Hamadeh, GenMab <ul style="list-style-type: none"> <li>• <b>Leveraging the NCI Cancer Research Data Commons</b> David Patton, Center for Biomedical Informatics and Information Technology, NCI</li> <li>• <b>Machine Learning and the Use of AI, an update on INFORMED</b> Sean Khozin, FDA</li> <li>• <b>Data-driven Knowledge Graphs and AI for Biomarker Discovery and Patient Selection</b> Janusz Dutkowski, Data4Cure</li> <li>• <b>Machine-learning Systems and Discovery of Actionable Biomarkers for Clinical Development</b> Pratik Shah, MIT</li> <li>• <b>Use of RWD in Accelerating Biomarker Discovery</b> Shrujal Baxi, FlatIron Health</li> <li>• <b>Discussion/Panel</b></li> </ul>
3:50 – 4:00 pm	<b>Wrap-up and Adjourn</b>