| Monday, May 13 | | | |
|---------------------|-------------------|--|---|
| Start Time | End Time | All tutorials will take place in the Mellon Institute Building at CMU | |
| 7:30 AM | 5:30 PM | Registration - location Alumni Hall, Connelly Ballroom | |
| | | Customized workflows for the interactive visualization of large-scale biological networks: A Python-based framework | |
| 8:00 AM | 12:00 PM | Room 328 of the Mellon Institute at CMU Beyond normalization: Incorporating scale into high-throughput sequencing analysis | |
| 8:00 AM | | Room 348 of the Mellon Institute at CMU | |
| 12:00 PM | 1:00 PM | LUNCH ON OWN Student-Focused Session: Navigating In-Person Conferences | |
| 1:00 PM | 5:00 PM | Room 328 of the Mellon Institute at CMU | |
| | | Applications of Generative AI, LLMs, and Traditional Machine Learning for Personalized Medicine and Drug Discovery | |
| 1:00 PM | 5:00 PM | Room 348 of the Mellon Institute at CMU | |
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| Tuesday, May 14 | | | |
| Start Time | End Time | Clapp Hall Auditorium | Cathedral of Learning, G24 |
| 8:00 AM | 5:00 PM | Registration - location Alu | mni Hall, Connelly Ballroom |
| 8:45 AM | 9:00 AM | Conference Welcome | |
| 9:00 AM | 10:00 AM | Keynote - Oana Carja | |
| 10:00 AM | 10:30 AM | | Coffee Break - Area TBD |
| 10:30 AM | 12:00 PM | General Track | Responsible AI for Multimodality Biomedical Data Analysis |
| 12:00 PM | 1:30 PM | | Lunch with Posters |
| 1:30 PM | | Keynote - Mingfu Shao | |
| 2:30 PM | | General Track | Responsible AI for Multimodality Biomedical Data Analysis |
| 3:30 PM | | Contract Track | Coffee Break - Area TBD |
| 4:00 PM | | General Track | Responsible Al for Multimodality Biomedical Data Analysis |
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| 6:00 PM | + | Bio Break | Responsible AI for Multimodality Biomedical Data Analysis |
| 6:15 PM | 7:30 PM | Education Session | responsible Afron Multimodality Biomedical Data Analysis |
| Made and the Africa | | | |
| Chart Times | Wednesday, May 15 | | |
| Start Time | End Time | Clapp Hall Auditorium | Cathedral of Learning, G24 |
| 8:00 AM | | | mni Hall, Connelly Ballroom |
| 8:45 AM | | Welcome - Day 2 | |
| 9:00 AM | | Keynote - Nathan Clark | |
| 10:00 AM | | | Coffee Break - Area TBD |
| 10:30 AM | | General Track | Computational Approaches for Spatial Omics |
| 12:00 PM | | | Lunch with Posters |
| 1:30 PM | 2:30 PM | Keynote - Sarah Bagby | |
| 2:30 PM | 3:30 PM | General Track | Computational Approaches for Spatial Omics |
| 3:30 PM | 4:00 PM | | Coffee Break - Area TBD |
| 4:00 PM | 6:00 PM | General Track | Computational Approaches for Spatial Omics |
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| | | Thursda | y, May 16 |
| Start Time | End Time | | Cathedral of Learning, G24 |
| 8:00 AM | | | mni Hall, Connelly Ballroom |
| 8:45 AM | 9:00 AM | Welcome - Day 3 | |
| 9:00 AM | 10:00 AM | Keynote - Daniel Lobo | |
| 10:00 AM | 10:30 AM | | Coffee Break - Area TBD |
| 10:30 AM | 12:00 PM | General Track | Scalable Analysis for Big Biological Data |
| 12:00 PM | 1:30 PM | | Lunch on Own |
| 1:30 PM | | General Track | Scalable Analysis for Big Biological Data |
| 3:30 PM | | | Coffee Break - Area TBD |
| 4:00 PM | | Keynote - Andreas Pfenni | |
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