

GALAXY PROTEOMICS @ ASMS 2015

June 1st 2015: Monday

WORKSHOP: The **Galaxy** Framework for MS-based Informatics.

Room 123/124 5:45pm - 7:00pm



June 2nd 2015: Tuesday

ORAL PRESENTATION: 9:50 AM ; Room 130/132 SearchGUI and PeptideShaker deployed in the **Galaxy** framework: A powerful informatics platform for protein identification and beyond. (Cooke *et al*)

- **POSTER 127:** Democratizing and expanding the reach of DIA Mass Spectrometry: Developing OpenSWATH tools and workflows within user-friendly **Galaxy-P** platform. (Jagtap *et al*)
- **POSTER 131:** Plugging Proteomics Peptide-Spectral Match Visualization into **Galaxy**. (Johnson *et al*)
- **POSTER 366:** Metaproteomic analysis of human cervical-vaginal fluid in residual Pap tests: Insights into the cervical microbiome. (Griffin *et al*)
- **POSTER 638:** Revealing Pathways In COPD-Associated Lung Cancer Large-Scale Quantitative Multiomic Analysis. (Sandri *et al*)

June 4th 2015: Thursday

- **POSTER 455:** A Novel Analytical-Informatics Platform Reveals the Hidden Tryptic Peptidome and Improves Multi-omic Applications. (Guerrero *et al*)