



Workshop

New mass spectrometry approaches to study cancer genome functions and beyond...

December 13th, 2018

Montpellier, IRCM Conference room

09:20 am	Welcome coffee
10:00 am	Session I Chairs: Jérôme Dejardin, Eric Julien
10:00 �	Deciphering chromatin biology using integrative omics approaches Michiel VERMEULEN, Radboud Institute of Molecular Life Sciences, Nijmegen, NL
10:40 �	The control of telomere maintenance by heterochromatin Jérôme Dejardin, Institute of Human Genetics, Montpellier
11:05�	Identification of replication factories components using mass spectrometry Cyril Ribeyre, Institute of Human Genetics, Montpellier
11:30 �	Exploring the biology of histone variants with mass spectrometry Jérôme GOVIN, Institute for Advanced Biosciences, La Tronche, France
12:10 pm	Lunch
14:00 pm	Session II Chairs: Kerstin Bystricky, Daniel Fisher
14:00 pm	Combining mass spectrometry with FACS for comprehensive proteomic and phosphoproteomic analysis of cell cycle transitions Tony LY, Centre for Gene regulation & Expression, University of Dundee
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14:00 �	Combining mass spectrometry with FACS for comprehensive proteomic and phosphoproteomic analysis of cell cycle transitions Tony LY, Centre for Gene regulation & Expression, University of Dundee A combined laser microdissection and proteomic analysis method for identification of robust biomarkers in oncology
14:00 * 14:40 *	Combining mass spectrometry with FACS for comprehensive proteomic and phosphoproteomic analysis of cell cycle transitions Tony LY, Centre for Gene regulation & Expression, University of Dundee A combined laser microdissection and proteomic analysis method for identification of robust biomarkers in oncology Anne-Aurélie Raymond, INSERM U1053, Liver cancer and invasion, Bordeaux Structural elucidation of cellular proteoforms by Top-down MS

