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| 09h15 - 09h55 | <b>Welcome Participants</b>                                                                                                                                                                |
| 09h55 - 10h00 | <b>WELCOME &amp; INTRODUCTION</b>                                                                                                                                                          |
| 10h00 - 11h00 | <b>Invited Speaker</b><br><i>Chair: Isabel Chillon, IGMM, Montpellier</i><br><br><b>Jorge FERRER</b> (CRG, Barcelona)<br><br><b>An RNA-Centric View of the Metabolic Gatekeeper HNF1A</b>  |
| 11h00 - 11h30 | <b>Coffee break</b>                                                                                                                                                                        |
|               | <b>SESSION 1 – Transcriptional and Post-Transcriptional Processes in the Nucleus</b><br><i>Chairs: Charlotte Grimaud, IGH, Montpellier &amp; Julie Carnesecchi, IGMM, Montpellier</i>      |
| 11h30 - 11h45 | <b>Flavia Mazzarda</b> (IGH, Montpellier)<br><i>Deciphering the inner workings of the HIV-1 promoter using MS2 imaging and SMF</i>                                                         |
| 11h45 - 12h00 | <b>Nazim Sarica</b> (IGH, Montpellier)<br><i>A multiplexed quantitative ddPCR approach to study hepatitis B virus transcriptional and post transcriptional regulation</i>                  |
| 12h00 - 12h15 | <b>Ruoyang Zhai, BGI, France</b><br><i>Enter the Multi-Omics Era with BGI Genomics: Technical Highlights</i>                                                                               |
| 12h15 - 12h30 | <b>Wan Xiang</b> (IGMM, Montpellier)<br><i>The RNA binding activity of the Drosophila Hox Transcription Factor Ultrabithorax contributes to muscle morphogenesis in developing embryos</i> |
| 12h30 - 12h45 | <b>Margot Lugoboni</b> (iGReD, Clermont-Ferrand)<br><i>Transcriptional regulation mechanisms mediated by m<sup>6</sup>A RNA modification in Drosophila melanogaster</i>                    |
| 12h45 – 14h30 | <b>LUNCH &amp; POSTER SESSION</b>                                                                                                                                                          |
|               | <b>SESSION 2 – From RNA Biogenesis to Translation</b><br><i>Chairs: David Lleres, CBS, Montpellier &amp; Anne-Catherine Prats, I2MC, Toulouse</i>                                          |
| 14h30 - 14h45 | <b>Chloé Bessiere</b> (CRCT, Toulouse)<br><i>Long-read sequencing identifies full-length circular RNAs linked with therapy resistance in pediatric cancers</i>                             |
| 14h45 - 15h00 | <b>Julia Schaeffer</b> (UGA, Grenoble)<br><i>An issue of translation: gene regulation mechanisms to unlock neuron regeneration</i>                                                         |
| 15h00 - 15h15 | <b>Javier Florido</b> (IRCM, Montpellier)<br><i>Understanding the epitranscriptomic landscape regulated by the p53 pathway</i>                                                             |
| 15h15 - 15h30 | <b>Léo Le Tareau</b> (CRBM, Montpellier)<br><i>A lncRNA-RNase mechanism at the center of the elimination process of nucleolus-related inclusions</i>                                       |
| 15h30 - 16h00 | <b>Break</b>                                                                                                                                                                               |
|               | <b>SESSION 3 – RNA Biology and Infection</b><br><i>Chairs: Flavia Mazzarda, IGH, Montpellier &amp; Yves Henry, CBI, Toulouse</i>                                                           |
| 16h00 - 16h15 | <b>Damien Avinens</b> (IRIM, Montpellier)<br><i>Deciphering the translation dynamics of SARS-CoV-2 UTRs</i>                                                                                |
| 16h15 - 16h30 | <b>Beata Jady</b> (MCD, Toulouse)<br><i>Epstein-Barr virus v-snoRNA1 modulates translation of the infected host cell</i>                                                                   |
| 16h30 - 16h45 | <b>Kuldeep Lahry</b> (IRCM, Montpellier)<br><i>Microbial metabolite preQ1 regulates host cell physiology through selective tRNA degradation and translation regulation</i>                 |
| 16h45 - 17h00 | <b>Johannes Keisers</b> (CBS, Montpellier)<br><i>mRNA degradation as a rescue mechanism for translation-inhibiting antibiotics</i>                                                         |
| 17h05 - 17h15 | <b>Presentation of Awards and Conclusion</b>                                                                                                                                               |
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