Schedule

9:30 - 10:15 **Guillaume Filion** (Genome Architecture Group Leader - CRG) *Chromatin plasticity*

10:15 - 10:40 **Neus Martínez** (Multicellular Systems Biology group) *Morphometrics showing genetics: semiautomatic tools for quantitative shape analysis in biomedical research*

10:40 - 11:05 **Felix Campelo** (Intracellular Compartmentation group) *How can lipids and proteins shape membranes to regulate protein export?*

11:05 - 11:30 Coffee break

11:30 - 11:55 **Lluís Morey** (Epigenetic Events in Cancer group) What are the molecular mechanisms that drive embryonic stem cell differentiation?

11:55 - 12:20 **Thomas Pengo** (Design of Biological Systems group) *Image analysis for quantitative biology at the nanoscale*

12:20 - 12:45 **Ombretta Foresti** (Organelle biogenesis and homeostasis) Sterol homeostasis requires regulated degradation of squalene monooxygenase by the ubiquitin ligase Doa10/Teb4

12:45 - 13:10 **Dmitry Pervouchine** (Computational Biology of RNA Processing group) *Conserved architecture of human and mouse transcriptomes*

13:30 - 14:30 Lunch

14:30 - 14:55 **Laurence Wurth** (Regulation of Protein Synthesis in Eukaryotes group) *Oncogenic properties of the RNA-binding protein UNR:Targets in melanoma*

14:55 - 15:20 **Juanjo Fraire** (Biomechanics of Morphogenesis / Comparative Analysis of Developmental Systems groups) *Biophysics and evolution - comparative tissue dynamics in flies and midges*

15:20 - 15:45 **Yolanda Schaerli** (Gene Network Engineering group) *A design space for synthetic stripe-forming networks*

15:45 - 16:15 Coffee break

16:15 - 16:40 **Daniela Sanges** (Reprogramming and Regeneration group) *Wnt/β-catenin signalling triggers neuron reprogramming and regeneration in the mouse retina*

16:40 - 17:40 **Joan Richtsmeier** (Invited speaker: Pennsylvania State University) *Opportunities for growth disguised as unsolvable problems*

17:40 - 18:00 Conclusion and closing remarks