

CRUK CoL Centre Cancer Evolution Symposium



9.15 - 9.45am	Registration	
9.45-10.00	Francesca Ciccarelli	Wellcome
Session 1	Chair: Magnus Haughey, QMUL	
10.00 am - 10.25am	Heba Sailem , KCL	Deep learning approaches for decoding tumour microenvironment heterogeneity
10.25am - 10.50am	Simone Zaccaria, UCL	Measuring proliferation rates of distinct clones in metastatic tumours using single-cell whole-genome sequencing
10.50am - 11.15am	Mirjana Efremova, QMUL	Dissecting cell heterogeneity and phenotypic plasticity in colorectal cancer metastasis
11.15am - 11.40am	Coffee break	
Session 2	Chair: Piyali Ganguli, Crick	
11.40am - 12.05pm	Xiao Qin, UCL	A Single-cell Perturbation Landscape of Colonic Stem Cell Polarisation
12.05pm - 12.30pm	Sheeba Irshad, KCL	TBA
12.30pm - 12.55pm	Maise Al Bakir, Crick	The evolution of metastases in non-small cell lung cancer
12.55pm - 2pm	Lunch	
Session 3	Chair: Jelmar Quist, KCL	
2.00pm - 2.25pm	Mariia Yuneva, Crick	Using mass spectrometry imaging to dissect metabolic heterogeneity of tumours
2.25 pm - 2.50pm	Lucia Montorsi, KCL	RNAscope coupled with Imaging Mass Cytometry allows high resolution validation of spatial transcriptomics
2.50pm - 3.15pm	Roberto Bellelli, QMUL	DNA Polymerase Epsilon and its roles in genome stability and mediating sensitivity to anti-cancer therapeutics
3.15pm - 4.00pm		Coffee break
Facilities	Chair: Carlos Martinez-Ruiz, UCL	
4.00pm - 4.20pm	Tristan Clark	The CoLCC computational hub
4.20pm - 4.40pm	Sam Marguerat	Single cell genomics @CoLCC
4.40pm - 5.00pm	Joseph Hartlebury	Tissue cyclic fluorescence (T-CyCIF) technology
5.00pm - 5.15pm	Nicky McGranahan	Conclusions

Networking