

Nº	TITLE	TUTORS	9:20 10:00	10:00 10:40	Coffee Break	11:00 11:40	11:40 12:20	12:20 13:00	Lunch	15:00 15:40	15:40 16:30	16:30 17:10
D01	The secret life of your crystallization drop: Do you know what really happens in your drops?	Bernhard Rupp	■		Coffee Break		■		Lunch		■	
D02	Characterization, analysis and use of crystallization screening outcome	Edward H. Snell		■		■		■				
D03	AlphaFold and biochemical considerations for protein crystallization	Joe Ng	■				■			■		
D04	Microfluidic tools and crystallization: from fabrication to application	Isaac Rodríguez		■				■				■
D05	Crystallization and crystallography in microchips	Claude Sauter	■				■				■	
D06	Gels for Crystallization	Ángeles Hernández		■		■		■				
D07	Protein crystallization by capillary counterdiffusion	Luis González	■				■			■		
D08	Hanging drop setup – you probably think you know how to do this already	Janet Newman		■		■		■				■
D09	How to perform vapour diffusion experiments: Hanging and Sitting drops	Ivana K.S. / Paulina R	■				■				■	
D10	The Temperature-controlled Optimization of Crystallization	M. Budayova-Spano		■				■		■		■
D11	Microbatch crystallization under oil	Lata Govada	■				■				■	
D12	Dialysis, crystallization and the Hofmeister series	Jeroen Mesters		■		■				■		
D13	Crystallization and Structural Studies on Protein-Nucleic Acid Complexes	C. Biertümpfel	■				■				■	
D14	Cryo-Electron tomography – method to bridge scales in biological system	Naoko Mizuno		■				■		■		■
D15	Growth of microcrystals of Photosystem I for serial femtosecond crystallography	P. Fromme / J.M. García / J-H. Yang	■			■					■	
D16	Labelling and Low-Cost Imaging of Macromolecule Crystals	M. Pusey / C. Tarver		■			■			■		
D17	Crystallization of membrane proteins in lipid mesophases	Martin Caffrey	■			■					■	
D18	The European X-Ray Free-Electron Laser Facility (EuXFEL)	Diogo Melo / Faisal H. Mekky Koua		■				■		■		■
D19	Microseed Matrix Screening experiments using a robot	S. Kolek / M. Sharpe	■				■				■	
D20	Formulatrix µPulse TFF - for Sample Concentration - Diafiltration (Buffer Exchange, Desalting)	Kate Lewis		■				■		■		
D21	The most sample efficient sample qualification method. Automated DLS	Karsten Dierks	■			■					■	
D22	Chromatography Principles and ÄKTA™ chromatography systems	P. Braga / E. García		■			■			■		
D23	Crystal fishing and mounting	Simon Tanley	■					■			■	