

	Friday Jun 2	Saturday Jun 3 <b>General Techniques</b>	Sunday Jun 4 <b>Biophysics and binding</b>	Monday Jun 5 <b>Informatics</b>	Tuesday Jun 6 <b>Small molecule chemistry</b>	Wednesday Jun 7	Thursday Jun 8 <b>Biologics</b>	Friday Jun 9 <b>Case Studies</b>	Saturday Jun 10	
8:45		Introduction (Directors & IT)								
9	ARRIVALS (Dinner provided)	Historical Perspective ( <b>T. Blundell</b> )	Measuring binding affinities ( <b>F. von Delft</b> )	Machine learning in drug discovery ( <b>C. Deane</b> )	General MedChem ( <b>M. Congreve</b> )	Conformational landscaping of the SARS-COV-2 spike protein ( <b>G. Verkhivker</b> )	Intro to Biologics ( <b>C. Deane</b> )	Structure-based drug discovery targeting HIV reverse transcriptase ( <b>E. Arnold</b> )	DEPARTURES	
9:45		Introduction to crystallography ( <b>C. Lesburg</b> )	Protein Conformational Landscapes ( <b>M. Fischer</b> )	Data integration & pharmacology ( <b>A. Bradley</b> )	Compound design ( <b>A. Bradley</b> )	Accelerating drug discovery ( <b>A. Kotecha</b> )	Rational design ( <b>S.J. Fleishman</b> )	Case study 3 ( <b>S. Jacob</b> )		
10:30		<i>Coffee</i>	<i>Coffee</i>	<i>Coffee</i>	<i>Coffee</i>		<i>Coffee</i>	<i>Coffee</i>		
11:00		CryoEM in drug discovery ( <b>G. Scapin</b> )	Computing Drug Binding Kinetics ( <b>R. Wade</b> )	Docking, Free energy calculation and Molecular Dynamics ( <b>S. Riniker</b> )	In Silico SB optimization for accelerated drug discovery ( <b>E. Davis</b> )	Talks from abstracts: G.A. Bezerra (4) J.C. Aschenbrenner (5)	Biologics Developability ( <b>R. Buick</b> )	Talks from abstracts: E. S. Cunha (13) M. Ruzskowski (45) S.S.R. Burman (44)		
11:45		High-throughput and automation ( <b>F. von Delft</b> )	Targeted Allosteric Modulators ( <b>G. Verkhivker</b> )	B. Vertessy (FEBS) Intro to FEBS	3D structures, interactions, and implications ( <b>AY Sheikh</b> )	D. Vattovaz (58) C. Catalano (10) A. Chopra (12)	Atoms up protein engineering ( <b>J. Williams</b> )	Z. Kozicka (27) V. Yadrykhins'ky (62)		
12:30		<i>Lunch</i>	<i>Lunch with poster preview</i>	<i>Lunch</i>		Excursion	<i>Lunch with poster preview</i>	<i>Lunch</i>		
2:30		Project Moonshot <b>A. VonDelft</b>	WS	WS	WS		WS	WS		Fragment-based design of MAT2 inhibitors ( <b>M. Schimpl</b> )
3:15		Machine learning and protein structure prediction: ( <b>C. Outeiral Rubiera</b> )								Cu' havi 'na bona vigna, havi pani, vinu e ligna" – Startups ( <b>J. Williams</b> )
4:00		<i>Coffee</i>	<i>coffee</i>	<i>Coffee</i>	<i>coffee</i>			<i>coffee</i>		<i>coffee</i>
4:30		WS Intro	WS	WS	WS		WS	WS		Where will SB be in 10 years? ( <b>S. Jacob</b> ) + group discussion
5:15	Ice Breaker	Ice breaker presentations								
6:00	Intro to Erice	Poster Session Odd Numbers								
8:00	Welcome buffet	Dinner at posters					Dinner at posters	Farewell dinner		

## Workshop schedule

Sunday – 4 June		Monday – 5 June		Tuesday – 6 June		Thursday – 8 June	
2:30-4:00	4:30-6:00	2:30-4:00	4:30-6:00	2:30-4:00	4:30-6:00	2:30-4:00	4:30-6:00
<b>Ligand crystallography /Hands on compound building</b> <i>C. Lesburg</i>	<b>Ligand crystallography /Hands on compound building</b> <i>C. Lesburg</i>	<b>General Machine Learning</b> <i>F. Boyles and C. O. Rubiera</i>	<b>General Machine Learning</b> <i>F. Boyles and C. O. Rubiera</i>	<b>Biologics design – Rosetta</b> <i>S. Fleishman</i>		<b>Hands-on single-particle cryoEM data analysis with cryoEDU</b> <i>M. Herzik</i>	
<b>General Machine Learning</b> <i>F. Boyles and C. O. Rubiera</i>	<b>General Machine Learning</b> <i>F. Boyles and C. O. Rubiera</i>	<b>Virtual Screening at Giga Scale</b> <i>S. Muchmore and M. Krier (OpenEye)</i>	<b>Virtual Screening at Giga Scale</b> <i>S. Muchmore and M. Krier (OpenEye)</i>	<b>In silico SB optimization for Accelerated Drug Discovery</b> <i>E. Abualrous and Z. Si Chaib (Schroedinger)</i>		<b>In silico SB optimization for Accelerated Drug Discovery</b> <i>E. Abualrous and Z. Si Chaib (Schroedinger)</i>	
<b>Hands-on single-particle cryoEM data analysis with cryoEDU</b> <i>M. Herzik</i>	<b>Hands-on single-particle cryoEM data analysis with cryoEDU</b> <i>M. Herzik</i>	<b>Drug Design using CSD-Discovery</b> <i>R. Chikhale and F. Montisci (CCDC)</i>	<b>Drug Design using CSD-Discovery</b> <i>R. Chikhale and F. Montisci (CCDC)</i>	<b>SmartEPU and G2</b> <i>A. Kotecha and A. Koh(TFS)</i>	<b>SmartEPU and G2</b> <i>A. Kotecha and A. Koh (TFS)</i>	<b>Biologics design – SabDab</b> <i>F. Boyles and C. O. Rubiera</i>	<b>Biologics design – SabDab</b> <i>F. Boyles and C. O. Rubiera</i>