S-SAD diffraction data phasing of macromolecule single crystals from home and Synchrotron X-ray sources Oeiras, October 4-6, 2007

October 4

8.45	Bus departs from Hotel Praia-Mar	
9.15	Anomalous Scattering in Macromolecular Crystallography	Gordon Leonard
10.50	Coffee break	
11.15	Introduction to strategies for in-house phasing using S-SAD and halide soaks.	George Sheldrick
13.00	Lunch	
14.30	Data collection with the X8 Proteum Diffractometer - Group 1 / Group2	Local organization
16.30	Introduction to the tutorial computing environment	Local organization
18h45	Bus departs from ITQB	

October 5

8.45	Bus departs from Hotel Praia-Mar	
9.15	Why we should make more use of softer X-rays in the de novo phasing of the crystal structures of macromolecules	Gordon Leonard
10.15	Coffee break	
10.30	On the use of Longer X-ray Wavelengths in Macromolecular Crystallography	Manfred Weiss
11.30	Scaling in-house data with SADABS	George Sheldrick
13.00	Lunch	
14.00	Scaling strategy for diffraction data from soft X-ray diffraction	Manfred Weiss
16.00	XPREP, SHELXC and hkl2map	George Sheldrick
17.30	Training with collected data	
18h45	Bus departs from ITQB	

October 6

8.45	Bus departs from Hotel Praia-Mar	
9.15	Locating substructures with SHELXD	George Sheldrick
10.15	Coffee break	
10.30	Phasing and density modification	Clemens Vonhrein
12.00	Burning questions	All Speakers
13.00	Lunch	
14.30	Density modification with SHELXE	George Sheldrick
16.00	SHARP	Clemens Vonhrein
17.30	Training with collected data	
18h45	Bus departs from ITQB	