



# **Intensive Training Course:**

# **SAXS and Computational Techniques to Study Intrinsically Disordered Proteins**

# EMBL, Hamburg, Germany, March 11 – 16, 2013

# **Program**

## Monday, 11th March

10:00 - 12:30	Arrival
12:30 - 14:30	Lunch
14:30 - 15:00	Welcome address
15:00 - 16:00	Introduction to SAXS (Dmitri Svergun)
16:00 - 16:30	Coffee break
16:30 - 17:30	Sample preparation (Rob Meijers)
17:30 - 18:30	Introduction to P12 beamline (Clément Blanchet)
18:30 – 20:00	BBQ

#### Tuesday, 12th March

09:00 - 10:30	Initial data processing (Al Kikhney)
10:30 - 11:00	Coffee break
11:00 - 12:30	Ab initio methods (Daniel Franke)
12:30 - 14:00	Lunch
14:00 - 15:30	Atomic structure based modeling (Maxim Petoukhov)
15:30 - 19:30	Measurements at the Beamline/SAXS data analysis practical

## Wednesday, 13th March

09:30 - 10:30	Mixture analysis (Peter Konarev)
10:30 - 11:00	Coffee break
11:00 – 12:30	Ensemble Optimization Method (EOM) application (Giancarlo Tria)
12:30 - 14:30	Lunch
14:30 – 17:30	Students' presentations:
14:30 – 14:50	Biao Fu
14:50 – 15:10	Cesyen Cedeno
15:10 – 15:30	Eduardo Calcada
15:30 – 15:50	Erica Valentini
15:50 – 16:10	Sergio Gil Caballero
16:10 - 16:30	Magdalena Korsak
16:30 - 16:50	Mikhail Kachala
16:50 – 17:10	Pallab Bhowmick
17:10 – 17:30	Priyanka Joshi
17:30 – 19:00	Supervisory board meeting

## Thursday, 14th March

09:30 - 11:00	Incorporation of SAXS measurements as structural restraints in molecular dynamics
	simulations (Michele Vendruscolo)
11:00 - 11:30	Coffee break
11:30 - 12:30	Round table discussion with IDPbyNMR Marie Curie fellows
12:30 - 14:30	Lunch
14:30 - 16:00	Students' presentations:
14:30 - 14:50	Jaka Kragelj
14:50 - 15:10	Tomas Hosek
15:10 - 15:30	Zsofia Solyom
15:30 – 15:50	Hadas Raveh-Amit
17:30 - 19:30	Measurements at the Beamline/SAXS data analysis practical

# Friday, 15th March

09:30 - 10:30	NMR and small angle scattering - tools for the characterisation of highly flexible systems ( <i>Martin Blackledge</i> )
10:30 - 11:00	Coffee break
11:30 – 12:30	Flexible-meccano: a tool for the generation of explicit ensemble descriptions of intrinsically disordered proteins and their associated experimental observables (Martin Blackledge)
12:30 - 14:30	Lunch
14:30 - 19:00	SAXS data analysis practical

# Saturday, 16th March

09:30 - 10:30	Discussion of obtained results
10:30 - 11:00	Coffee break
11:00 – 12:30	Discussion of obtained results
12:30 - 14:30	Lunch
14:30	Departure