

Biomathematics afternoon in Ljubljana, 2016

Wednesday, 17 February 2016

First session - 2.05 (15:00 - 16:20)

time	[id] title	presenter
15:00	[0] Bernhard Thiel (TBI) - Integrating Cryo-EM with Coarse Grained RNA 3D Structure Prediction in ERNWIN	
15:20	[1] Roman Jerala (KI) - Design principles for rapid folding of knotted DNA	
15:40	[2] Ronny Lorenz (TBI) Constraints in RNA secondary structure prediction	
16:00	[3] Ajasja Ljubetič (KI) - Designing, modelling and folding protein polyhedra	