

Programme for the Bioinformatic meeting: “Next generation sequencing” at Lian 2008!

09.00 – 09.15	Registration
09.15 – 09.30	Welcome by Finn Drabløs
09.30 – 10.15	Joost Martens , Nijmegen Center For Molecular Life Sciences (NCLMS), Department of Molecular Biology “High throughput sequencing for understanding the epigenetics of Leukemia”
10.15 – 11.00	Anne-Mette K. Hein, CLC Bio ”Some issues in the analysis of mRNA-seq data”
11.00 – 11.30	Fruit break / Coffebreak
11.30 – 12.15	Nils Peder Willassen, Dept. of Medical Biology, University of Tromsø "Next generation sequencing in bioprospecting - challenges and opportunities"
12.15 – 12.45	Jan Egil Afseth, Dept. of Medical Microbiology, St. Olavs “Use of genome sequencing for identification and characterization of virulence factors in <i>E. coli</i> , group B streptococci and <i>Legionella pneumophila</i> .”
12.45 – 14.00	Lunch
14.00 – 14.45	Lex Nederbragt, CEES, Dept. of Biology, University of Oslo "High-Throughput Sequencing Services at the University of Oslo"
14.45 – 15.30	Inge Jonassen, Department of Informatics, University of Bergen “Bioinformatics challenges in the cod genome project”
15.30 – 16.00	Fruit break / Coffebreak
16.00 – 16.30	Mimmi Throne-Holst, SINTEF Materialer og kjemi "High Throughput Sequencing of Metagenomic DNA from an Extreme Environment"
16.30 – 17.00	Per Winge, Dept. of Biology, NTNU “High throughput sequencing: challenges and opportunities in plant research”
17.00 – 17.30	Hans Stenøien , Museum of Natural History and Archaeology, NTNU "Biodiversity research using next generation sequencing technologies"
17.30 – ca 19.30	Dinner