

## **Program for the Bioinformatic meeting: “High throughput –omics!” at Ringve 2010!**

09.00 – 09.15	Registration
09.15 – 09.30	Welcome by Finn Drabløs
09.30 – 10.15	<b>Vessela Kristensen</b> , Department of Genetics, Oslo University Hospital “High throughput genomic scans using SNPs”
10.15 – 11.00	<b>Tone Bathen</b> , Department of circulation and medical imaging, NTNU “High throughput metabolomics using NMR”
11.00 – 11.30	Fruit break / Coffebreak
11.30 – 12.15	<b>Harald Barsnes</b> , The Department of Biomedicine, UiB “Bioinformatics and Quantitative Proteomics”
12.15 – 12.45	<b>Eiliv Lund</b> , Samfunnsmedisin, UiT “High throughput expression analyses in systems epidemiology – challenges and experiences from NOWAC”
12.45 – 14.00	Lunch
14.00 – 14.45	<b>Dag Coucheron</b> , UiT ““
14.45 – 15.15	<b>Åsa Johansson</b> , Uppsala Clinical Research Center, Uppsala University Hospital, Sweden “Genome-wide association analyses and whole-genome transcription profiling”
15.15 – 15.45	Fruit break / Coffebreak
15.45 – 16.15	<b>Trevor Clancy</b> , Pubgene “Coremine Medical for analysis of immune network signatures in gene expression and survival”.”
16.15 – 17.00	<b>Eivind Almaas</b> , Department of Biotechnology, NTNU “Gene-expression networks in human and chimp brains”
17.00 – 17.30	<b>Per Bruheim</b> , Department of Biotechnology, NTNU “MS based Metabolomics: current status and future challenges”
17.30 – ca 19.00	Dinner