

# NTNU Bioinformatics Network Meeting

## Exomatics - Bioinformatics challenges of exome analysis

November 8th 2013

Suhm-huset - Øya – Trondheim

- 08.45-09.00 Registration
- 09.00-09.15 Welcome
- 09.15-09.45 Einar Ryeng – Genomics Core Facility – NTNU  
*Exome sequencing*
- 09.45-10.15 Sigve Nakken - Dept. of Tumor Biology – Oslo University Hospital  
*TBD*
- 10.15-10.45 Break (coffee,fruit)
- 10.45-11.15 Eivind Coward (Norwegian Cancer Genome Consortium, Trondheim)  
*Bioinformatics for the multiple myeloma exome project*
- 11.15-11.45 Gard Thomassen - Research Support Services Group – UiO  
*Services for sensitive data*
- 11.45-12.45 Lunch
- 12.45-13.15 Håkon Gjessing - Department of Genes and Environment - Norwegian Institute of Public Health  
*Family trio versus case-control designs in genetic association studies*
- 13.15-13.45 Oddgeir Lingaas Holmen - The Nord-Trøndelag health study  
*Exome arrays - a cost effective approach in population genetics*
- 13.45-14.15 Break (coffee)
- 14.15-14.45 Henrik Jensen – Centre for Biodiversity Dynamics - NTNU  
*Using genomic tools to understand eco-evolutionary dynamics of natural bird populations*
- 14.45-15.00 Closure
- 15.00-18.00 Dinner at Suhm-huset

Registration at: <http://tare.medisin.ntnu.no/seminar2013/registrate.php>