

## Program for DSKMs virus, metagenomics and WGS meeting.

**Venue:** Herlev Hospital, Lille auditorium ved hovedindgangen.

**Date:** Monday June 12, 2017. 10:00 to 15:00

Tilmelding til [Hanne.Lykke.Pedersen@regionh.dk](mailto:Hanne.Lykke.Pedersen@regionh.dk) senest 2. juni. Af hensyn til bestilling af frokost

### Program

- 9.30-10.00 Kaffe og registrering
- 10.00-10.40 Ed Feil, University of Bath, UK. The promises of whole genome pathogen sequencing.
- 10.40-11.20 Martin Sikora, Centre for GeoGenetics, University of Copenhagen. Microbial DNA in archaeological samples. Can we find DNA viruses?
- 11.20-12.00 Jens Bukh, Hvidovre Hospital. Role of NGS in HCV treatment.
- 12-13 Lunch on the Balcony with Networking
- 13.00-13.30 Anders Fomsgaard, SSI. Use of fast Micro-array metagenomic diagnostic of all virus in clinical samples.
- 13.30- 13.50 Søren Karst, Center for Microbial Communities, Aarhus. Using the Naopore MinION for metagenomics analysis.
- 13.50-14.10 Kim Ng, SSI. SSI WGS surveillance QC.
- 14.10-14.30 Roland Thele and Thomas Arn Hansen. A clinical sample negative in all microbial analysis but positive by microbiome sequencing.
- 14.30-14.50 Peder Worning. Dru-repeats and the origin of the staphylococcal *SCCmec* cassette.
- 14.50 Henrik Westh, Hvidovre Hospital. This is the end