

CRG Core Facility Technology Symposium

PROGRAM

“Proteomics 2.0.¹⁰”

5th February 2010

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|---------------|---|
| 10:00 – 10:15 | Welcome |
| 10:15 – 11:00 | Jakob Bunkenborg , Institute for Biochemistry & Molecular Biology, University of Southern Denmark, Denmark

“The landscape of post-translational modifications seen through a mass spectrometer” |
| 11:00 – 11:45 | Kris Gevaert , VIB Dept. of Medical Protein Research, UGent Proteome Analysis & Bioinformatics Unit, Belgium

“The Ends of Proteins...” |
| 11:45 – 12:15 | Coffee Break |
| 12:15 – 13:00 | Bernard Delanghe , Thermo Fisher Scientific, Bremen, Germany

“A new automated tool for SILAC quantitation” |
| 13:00 – 13:45 | Robert J. Beynon , Proteomics & Functional Genomics Group, Faculty of Veterinary Science, University of Liverpool, UK

“QconCATs – what’s the CATch?” |

