

Time	Event	Venue
11:00-11:15	Welcoming tea and coffee	BLT2
11:15-12:15	<i>Segmentation with Hidden Markov Models</i> . Dr Jüri Lember, Institute of Mathematical Statistics, Tartu University (Estonia)	BLT2
12:30-1:30	Lunch break (lunch provided free of charge for the invited speakers only, but everyone is welcome to join the group)	SCR (Founders building)
2:00-3:00	<i>Decision theoretic approaches to segmentation problems in genomics using Hidden Markov Models</i> . Dr Christopher Yau, Department of Mathematics, Imperial College London	BLT2
3:00-3:30	Discussion over tea and coffee	BLT2
3:30-4:30	Nonparametric testing of conditional independence. Dr Wicher Bergsma, Department of Statistics, LSE	BLT2
4:30-5:00	Closing discussions	BLT2

Abstracts of the invited talks:

1. *Segmentation with Hidden Markov Models*. Dr Jüri Lember, Institute of Mathematical Statistics, Tartu University (Estonia).
Abstract: We consider a unified approach to the segmentation/decoding problem based on the Bayesian decision theory. Our focus is on using Hidden Markov Models, where this approach allows us to re-examine various properties of some of the well-known decoders as well as develop a new class of path inference methods. Asymptotic properties of several of such segmentation methods will also be discussed.
2. *Decision theoretic approaches to segmentation problems in genomics using Hidden Markov Models*. Dr Christopher Yau, Department of Mathematics, Imperial College London
Abstract: Signal segmentation is a fundamental ingredient in many bioinformatics analyses. Hidden Markov Model (HMM) based approaches are a popular statistical model for these problems due to their simplicity and computational tractability. In this talk, I will discuss some applications of HMMs in cancer genomics and the application of Bayesian decision theoretic ideas to obtain flexible summary statistics.
3. *Nonparametric testing of conditional independence*. Dr Wicher Bergsma, Department of Statistics, LSE
Abstract: Random variables Y and Z are conditionally independent given X if, knowing the value of X , information about Y does not provide information about Z . Conditional independence relations are the building blocks of graphical models, applications of which include information extraction, speech recognition, computer vision, decoding of low-density parity-check codes, modeling of gene regulatory networks, gene finding and diagnosis of diseases, and graphical models for protein structure. The present talk discusses a new method to test conditional independence. Existing literature on the topic is usually restricted to the normal and categorical cases, but recently nonparametric testing has also received a fair amount of attention. Our method is also nonparametric, but differs from previous ones in that it is based on the following decomposition, which gives some advantages: Denote by $\Psi_{g,h}(x)$ the conditional covariance between $g(Y)$ and $h(Z)$ given $X=x$. Conditional independence of Y and Z given X holds if and only if the following two conditions hold: 1) For arbitrary g and h , $E[\Psi_{g,h}(X)] = 0$ 2) For arbitrary g and h , $\Psi_{g,h}(x)$ does not depend on x . Each condition can be tested separately. However, there are some technical difficulties which we explain and for which we provide a solution.

Organizers: Dr Alexey Koloydenko email: Alexey.Koloydenko@RHUL.ac.uk
Dr Teo Sharia email: T.Sharia@RHUL.ac.uk