

**13 MARCH 2017**

# **TB IS CHANGING**

**ST CATHERINE'S COLLEGE  
OXFORD**

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From 2017 all TB samples in England will have routine whole genome sequencing performed. This one-day conference will be an essential update for practitioners on the upcoming launch of routine whole genome sequencing and the implications for TB diagnosis, management and control.

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**For programme details and ticketing visit**

**<https://tbischanging.eventbrite.co.uk/>**

**£45 (£65 AFTER 23 JAN 2017) – CONCESSION**

**£75 (£95 AFTER 23 JAN 2017) – STANDARD**

**including lunch and wine reception**

# PROGRAMME

<b>Derrick Crook</b> PHE	<i>Introduction</i>
<b>Ibrahim Abubakar</b> UCL <b>Peter White</b> PHE	<i>The changing epidemiology of tuberculosis in England</i>
<b>Grace Smith</b> PHE	<i>The changing structure of TB services in England</i>
<b>Ana Cruz</b> University of Oxford	<i>Changes in phenotypic drug sensitivity testing</i>
<b>Tim Walker</b> University of Oxford	<i>Predicting drug susceptibility from Whole Genome Sequencing</i>
<b>David Wyllie</b> University of Oxford	<i>Whole Genome Sequencing compared to MIRU-VNTR for outbreak detection</i>
<b>Tim Peto</b> University of Oxford	<i>New format of TB reference laboratory reports</i>
<b>Gerry Davies</b> University of Liverpool	<i>Round-up of recent clinical trials and their implications</i>
<b>Graham Bothamley</b> N.E. London TB Services	<i>New drugs – access and experience of use</i>
<b>Marc Lipman</b> UCL/ Royal Free Hospital	<i>The impact of new laboratory data on clinical practice</i>
<b>Onn Min Kon</b> Imperial College NHS Trust	<i>Future of MDR-expert advice services</i>
<b>Sarah Anderson</b> PHE	<i>The role of TB control boards in local TB prevention and control</i>
<b>Martin Dedicoat</b> Birmingham Heartlands	<i>Early experiences of using WGS data for outbreak detection in Birmingham</i>
<b>Lauren Ahyow</b> PHE	<i>Role for social network analysis - fewer but more focused investigations?</i>
<b>Al Story</b> Find & Treat Service, UCLH	<i>Experience from the Find and Treat programme – a model to be replicated?</i>
<b>Hanif Esmail</b> University of Oxford	<i>Future of screening for latent TB</i>
<b>Kate Gaskell</b> LSHTM	<i>Developing an MDR-TB contact registry</i>
<b>Chris Conlon</b> Oxford University Hospitals <b>Helen Booth</b> UCLH <b>Martin Dedicoat</b> Birmingham Heartlands	<i>Case presentations and panel discussion</i>
<b>David Moore</b> LSHTM	<i>Closing remarks and round-up</i>