



Workshop

Genome-wide methods in cancer genetics
(Wykorzystanie technik analizy genomu w badaniach nowotworów)

28th October 2014,
Centrum Dydaktyczne Warszawskiego Uniwersytetu Medycznego, Aula B
ul. Księcia Trojdena 2a

AGENDA

9:00 - 9:15	Registration
9:15 - 9:30	Opening and foreword Prof. RAFAŁ PŁOSKI , Warsaw Medical University
9:30-10:15	<i>From custom genomic microarrays through target enrichment sequencing assays to identification of mutations predisposing to an inherited disorder: a case of multiple schwannomas</i> dr ARKADIUSZ PIOTROWSKI , Medical University of Gdańsk, Poland
10:15-10:30	Discussion
10:30-10:45	Coffee break
10:45-11:30	<i>Tracking tumor-specific changes from plasma</i> dr ELLEN HEITZER , Medical University in Graz, Austria
11:30-11:45	Discussion
11:45-12:30	<i>Single cell genomics - technologies and applications</i> ALEKSANDRA KOŁODZIEJCZYK , EMBL-European Bioinformatics Institute & Wellcome Trust Sanger Institute, Cambridge UK
12:30-12:45	Discussion
12:45-13:45	Lunch
13:45-14:15	<i>Making the switch in Molecular Pathology: from classical sequencing approach to NGS with IonTorrent, an opportunity for diagnostic and research,</i> dr LUCA QUAGLIATA , Institute of Pathology, University Hospital, Basel, Switzerland
14:15-14:30	Discussion
14:30-14:45	Coffee break
14:45-15:15	<i>Sensitive and accurate detection of mutations using Pacific Biosciences Circular Consensus Sequencing</i> dr ALEX GARVIN , Droplet Diagnostics, Mulhouse, France
15:15-15:30	Discussion
15:30	Closing the workshop

Further informations and applications:
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