

**THURSDAY 01.06.2017**

**9.00 – 9.15 REGISTRATION, COFFEE BREAK**

**9.15 -9.30 Presentation of STREAM project**  
**Magdalena Winiarska, PhD, Medical University of Warsaw, Poland**

**9.30 – 10.30 PrimeFlow RNA® : Cytometry-based analysis of RNA expression at single-cell level**  
**Marc Reudelsterz, Thermo Fisher Scientific**

**10.30 – 10.35 Discussion**

**10.35 – 10.55 Regulation of germinal center response by transcription factors using Prime Flow RNA method**  
**René Rauschmeier, Research Institute of Molecular Pathology, Vienna, Austria**

**10.55 – 11.00 Discussion**

**11.00 – 11.30 COFFEE BREAK**

**11.30– 12.10 Secrets of CST and standardization in flow cytometry**  
**Marzena Biernacka, BD, Poland**

**12.10– 12.15 Discussion**

**12.15 – 12.55 Applying panel design rules and optimizing panels for improved data quality**  
**Rafal Januszewski, BD, Poland**

**12.55 – 13.00 Discussion**

**13.00 – 14.00 LUNCH**

**14.00 – 14.40 Flow-cytometry based approaches to study the role of B-cell receptor in the pathogenesis of CLL**  
**Stefania Gobessi, PhD, ICGEB-Molecular Hematology Lab, Trieste, Italy**

**14.40 – 14.45 Discussion**

**14.45 – 15.25 Monitoring NK cell repertoire diversity and homeostasis by flow cytometry**  
**Aline Pfefferle, PhD, Oslo University Hospital, Norway**

**15.25 – 15.30 Discussion**

**15.30 – 16.00 COFFEE BREAK**