## **ICGEB Workshop on "Next Generation Diagnostics"**

22/03/2018-24/03/2018

Macedonian Academy of Sciences and Arts (MASA)
Skopje, Republic of Macedonia

12:45 - 14:15

Lunch



Venue: Macedonian Academy of Sciences and Arts

Bul. Krste Misirkov 2, Skopje, Republic of Macedonia

## **PROGRAM**

22 March 2018	
15:00 – 15:30	Welcome address and opening remarks
	Momir Polenakovic, Macedonian Academy of Sciences and Arts, Skopje
	<b>Taki Fiti,</b> President, Macedonian Academy of Sciences and Arts, Skopje
15:30 – 17:45	NGS diagnostics: State of the art and beyond
	(moderators: Richard Rosenquist and Dimitar Efremov)
	Clare Turnbull, Cambridge (Genomics England Initiative)
	Richard Rosenquist, Stockholm (Transferring NGS to clinical diagnostics)
	<b>Ulf Landegren</b> , Uppsala (New technologies for biomarker discovery)
17:45 – 18:00	Coffee break
18:00 - 19:15	NGS diagnostics in inherited diseases
	(moderators: Richard Rosenquist and Dimitar Efremov)
	Marcel Nelen, Nijmegen (Using NGS in a clinical diagnostic laboratory)
	Anna Wedell, Stockholm (Whole-genome sequencing in inherited diseases)
19:15 - 20:00	Welcome reception
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14:15 – 16:00	NGS diagnostics in hematological malignancies (moderators: Dimitar Efremov and Irina Panovska-Stavridis)
	Torsten Haferlach, Munich (NGS-based diagnostics of myeloid neoplasms)  Davide Rossi, Bellinzona (NGS-based diagnostics of lymphoid neoplasms)  Kostas Stamatopoulos, Thessaloniki (Immunogenetics in the era of NGS)
16:00 – 16:30	Coffee break
16:30 – 18:15	Predictive biomarkers and personalized treatment strategies (moderators: Richard Rosenquist and Momir Polenakovic)  Katerina Davalieva, Skopje (Proteomics of prostate cancer)  Päivi Östling, Stockholm (Precision medicine approaches in solid tumors)  Dimitar Efremov, Trieste (Targeting the kinome in B-cell malignancies)
18:15	Poster session

## 24 March 2018

24 March 2018	
	Bioinformatics analysis of NGS data
	(moderators: Fotis Psomopoulos and Valtteri Wirta)
09:15 - 09:45	Welcome and Introductions:
	- Outline of the general concept and particular content of the session.
	- Description of the informatics infrastructure and setup required for the tools and pipelines)
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09:45 – 10:15	<b>Valtteri Wirta,</b> Stockholm (Targeted Deep Sequencing: Implementing targeted sequencing for patient selection for clinical trials and for diagnostic purposes)
10:15 - 10:45	<b>Fotis Psomopoulos,</b> Thessaloniki (Targeted Deep Sequencing: End-to-end computational workflows towards the analysis of RNA-Seq data )
10:45 - 11:15	Henrik Stranneheim, Stockholm (Whole Exome / Whole Genome Sequencing:
	Informatics strategy for whole-genome sequencing based rare disease diagnostics)
11:15 - 11:45	Andreas Agathaggelidis, Thessaloniki (Whole Exome / Whole Genome
	Sequencing: Employing WGS approaches towards identifying key variants in CLL
	ontogenesis.
11:45 - 12:00	Discussion / Wrap-up:
	- General comments
	- Highlight key concepts that need to be taken into consideration for the

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- Discussion / Interaction with the participants