

Program of the 8th Microsymposium on Small RNAs

May 27th - 29th , 2013, IMBA, Vienna

Monday 27th

11.00 – 11.10 Welcome and Introduction

11.10 – 13.0	OStructure and Mechanism (1)
11.10 – 11.40	Elisa Izaurralde – Mechanisms of miRNA-mediate gene silencing
11.40 – 12.10	Jeremy Wilusz (Sharp Lab) – Changing dogmas of post-transcriptional control:
	Lessons from the MALAT1 noncoding RNA
12.10 – 12.30	Thermo Scientific – Tools for Analyzing MicroRNA Function in Mammalian Cells
12.30 – 13.00	Carolyn Phillips (Ruvkun Lab) - Compartmentalization of siRNA and piRNA
	pathways in C. elegans

13.00 – 14.00 Lunch

14.00 – 15.00 **Keynote Lecture**

Narry Kim - Regulation of microRNA through alternative processing and modifications

15.00 – 16.4	PhD Workshop – Part 1
15.00 – 15.20	Daniela Praher (Technau Lab, University of Vienna) – Cnidarian microRNAs regulate their targets by a plant-like cleavage mechanism
15.20 – 15.40	Alfred Bronkhorst (van Rij Lab, Nijmegen Centre for Molecular Life Sciences) – The DNA virus Invertebrate iridescent virus 6 is a target of the Drosophila RNAi machinery
15.40 – 16.00	Judith Hauptmann (Meister Lab, University of Regensburg) – Turning catalytically inactive human Argonaute proteins into active slicer enzymes
16.00 – 16.20	· · · · · · · · · · · · · · · · · · ·
16.20 – 16.40	Sophie Wöhrer (Mochizuki Lab, IMBA) – Assembly of a functional small RNA effector complex in Tetrahymena

16.40 – 17.10 Break

17.10 – 19.00	Genome detense (1)
17.10 – 17.40	Jean-René Huynh – tRNA fragments and transposable elements silencing in the
	Drosophila germline
17.40 – 18.10	Ramesh Pillai - Role of yet another tudor domain protein Tdrd12 in mouse piRNA
	biogenesis and its essential role for transposon silencing
18.10 – 18.30	Qiagen - To be determined
18.30 – 19.00	Tai Montgomery (Ruvkun Lab) – Crosstalk between small RNA pathways in C.
	elegans

19.00 – 19.20 EMBO Journal

19.30 Dinner at the IMBA/IMP cafeteria

Tuesday 28th

9.00 – 10.30	Small RNA Biology (1)
9.00 - 9.30	Richard Carthew – Regulation of miRISC activity by lipid signaling
9.30 - 10.00	Sue-Jean Hong (Bartel Lab) – Biological roles of miRNA-200 regulation of Zeb1 and
	Zeb2 during mammalian development
10.30 – 11.00	Antonio Giraldez – How life begins: The maternal to zygotic transition

10.30 - 11.00 Break

11.00 – 12.30 Genome Defense (2) 11.00 – 11.30 Shiv Grewal – Epigenetic genome control by RNAi and heterochromatin machinery 11.30 – 12.00 Kuniaki Saito – DmGTSF1 is required for effective retrotransposon silencing by Piwi and ovarian development in Drosophila 12.00 – 12.30 Eric Miska – Social RNA

12.30 - 13.30 Lunch

13.30 – 15.00	Structure and Mechanism (2)
13.30 – 14.00	Yukihide Tomari – Assembly and function of RISC
14.00 – 14.30	Illumina - ENCODE your research: Gene Expression and Regulation Update
14.30 – 15.00	Markus Hafner (Tuschl Lab) – RNA targets of LIN28

15.00 – 15.30 Coffee break / Poster viewing

15.30 – 16.20	Mechanism and Application
15.30 – 16.00 Anasta	sia Khvorova - RNA Therapeutics: Promise and Reality
16.00 - 16.20 Christo	f Fellmann (Mirimus) – A processing-optimized backbone boosts the
effective	eness of experimental miRNA-based RNAi

16.20 – 16.50 **Helen Pickersgill** – Scientific writing from an editor's perspective

17.00 - Tour, Dinner with academic speakers

Wednesday 29th

Ī	9.00 – 10.30	Small RNA Biology (2)
١	9.00 - 9.30	Chris Hammell – The coupling of temporal gene expression and cell fate specification
ı	9.30 - 10.00	Ana Eulalio – Shedding light on microRNA function through High Content Screening
ı	10.00 – 10.30	Petr Svoboda – The real RNAi – a tale of naughty mice

10.30 - 11.00 Break

11.00 – 13.00 PhD Workshop – Part 2 11.00 – 11.20 Felix Muerdter (Hannon Lab, CSHL)– A genome-wide RNAi screen draws a genetic framework for transposon control and primary piRNA biogenesis in Drosophila 11.20 – 11.40 Mirela Marasovic (Halic Lab, Gene Center Munich) – Argonaute and Trimmer generate Dicer-independent priRNAs and mature siRNAs to initiate heterochromatin formation 11.40 – 12.00 Zhao Zhang (Zamore Lab, UMass Medical School) – Rhino binding and convergent transcription are sufficient to generate a piRNA-producing locus 12.00 – 12.20 Matyas Ecsedi (Großhans Lab, FMI Basel)– Quantitative analysis reveals extensive target specificity of individual let-7 miRNA family members in vivo 12.20 – 12.40 Grzegorz Sienski (Brennecke Lab, IMBA) – Dissection of the Piwi-mediated transcriptional silencing process

13.00 – 15.00 Lunch and poster viewing

5.00 – 16.30 Structure and Mechanism (3)
15.00 – 15.30 Dinshaw Patel – Structural Biology of Prokaryotic and Eukaryotic Argonautes
15.30 – 16.00 Ryuya Fukunaga (Zamore Lab) – Regulation of Dicer activity by partner proteins and
a small molecule
16.00 – 16.30 Marina Chekulaeva – Exploring the mechanisms of miRNA function

16.30 – 17.00 Break

17.00 – 18.20	New Frontiers
17.00 – 17.30	Grzegorz Kudla – Mapping human miRNA interactions by CLASH
17.30 – 17.50	Exiqon - MicroRNA in biofluids - Robust biomarkers for disease
17.50 – 18.20	Uli Elling – Forward and reverse genetics in haploid embryonic stem cells

18.20 – 18.30 Award and closing of the meeting

18.30 - Dinner at IMBA