



## **CGC/CBG** meeting Epigenetics and non-coding RNAs

10 & 11 November 2011

Koninklijk Instituut voor de Tropen, Mauritskade 63, Amsterdam

## **Thursday November 10**

9:00- 9.30	Registration & welcome with coffee
	Chair Reuven Agami
9.30 – 10.00	<b>Witold Filipowicz</b> (Friedrich Miescher Institute, Basel, Switzerland) Mechanisms and regulation of miRNA-mediated repression in metazoa
10.00 – 10.30	<b>Robert Singer</b> (Albert Einstein College of Medicin, New York, NY, USA) Following single RNAs in living cells
10.30 – 11.15	Break
11.15 – 11.45	Ramin Shiekhattar (The Wistar Institute, Philadelphia, PA, USA) Elucidating the function of enhancer-like RNAs
11.45 – 12.30	Short talk by <b>Jarno Drost</b> (NKI-AVL, Amsterdam) p-53 dependent enhancer RNAs
	Short talk by <b>Selina van Leeuwen</b> (LUMC, Leiden) Involvement of long non-coding RNAs in regulating transcription of <i>Sox</i> genes
12.30 - 13.30	Lunch
	Chair Maarten van Lohuizen
13.30 – 14.00	<b>Kristian Helin</b> (Copenhagen Biocenter, Copenhagen, Denmark) Functional roles of TET proteins and hydroxymethylation in stem cells and cancer
14.00 – 14.30	<b>Bing Ren</b> (University of California, La Jolla, CA, USA) Base-resolution analyses of parent-of-origin and sequence dependent allele specific DNA methylation in the mouse genome
14.30 – 15.15	Break
15.15 – 16.00	Short talk by <b>Valerio Orlando</b> (Dulbecco Telethon Institute, Rome, Italy) A global role for chromatin associated RNAi components in transcriptional regulation
	Short talk by <b>Leonie Kamminga</b> (Hubrecht Institute, Utrecht) Ezh2 and the maintenance of tissue homeostasis during embryonic development of the zebrafish
16.00 – 16.45	<b>Eran Segal</b> (Weizmann Institute of Science, Rehovot, Israel) Unraveling the rules of transcriptional regulation using large-scale libraries of designed promoter sequences





## Friday November 11

	-
9.00 - 9.30	Registration & welcome with coffee
	Chair Bas van Steensel
9.30 – 10.00	<b>Edith Heard</b> (Institut Curie, Paris, France) Non-coding RNAs in X inactivation
10.00 – 10.30	Wolf Reik (London Research Institute, London, UK) Imprinting and non-coding RNAs
10.30 – 11.15	Break
11.15 – 11.45	<b>Mikiko Siomi</b> (Keio University School of Medicine, Tokyo, Japan) PIWI-interacting RNA in Drosophila germline
11.45 – 12.15	Klaus Rajewsky (Harvard Medical School, Boston, MA, USA) Hunting functional miRNA targets
12.15 – 13.30	Lunch
	Chair Thijn Brummelkamp
13.30 – 14.00	<b>Haruhiko Siomi</b> (Keio University School of Medicine, Tokyo, Japan) Maelstrom, a component of the piRNA pathway, coordinates microtubule organization during Drosophila oogenesis through interaction with components of the MTOC
14.00 – 14.30	<b>Robert Darnell</b> (The Rockefeller University School, New York, NY, USA) Maps of the RNA world
14.30 – 15.00	Break
15.00 – 15.30	<b>Howard Chang</b> (Stanford University, Stanford, CA, USA) Genome regulation by long noncoding RNAs
15.30 - 16.00	Nahum Sonenberg (McGill University, Montreal, Canada) Mechanism of action of miRNA
16.00	Closure