

Central European Institute of Technology BRNO | CZECH REPUBLIC

CEITEC: Building Towards Excellence Czech and Danish Scientists' Workshop on Life Sciences

Markus Dettenhofer

Executive Director

Copenhagen, Denmark 29 April 2014



EUROPEAN UNION EUROPEAN REGIONAL DEVELOPMENT FUND INVESTING IN YOUR FUTURE



OP Research and Development for Innovation





CEITEC is a scientific centre in the fields of life sciences, advanced materials and technologies whose aim is to establish itself as a recognized European center of science.





Vision Roadmap





Partnering institutions In Brno, Czech Republic



Masaryk University www.muni.cz



Brno University of Technology <u>www.vutbr.cz</u>



Mendel University in Brno

Mendel University in Brno www.mendelu.cz



University of Veterinary and Pharmaceutical Sciences Brno <u>www.vfu.cz</u>



Institute of Physics of Materials, Academy of Sciences of the Czech Republic www.ipm.cz



Veterinary Research Institute www.vri.cz



Management Structure



Measuring Success Against Global Standards



Board consists solely of Western European and US Members

Evaluation Sept. 2014 based on past performance, vision plan, and laboratory visit



Research Programmes



61 Group Leaders



CEITEC Publications in Q1%





Measures of Excellence EC definition

- critical mass of high level scientists
- integrating connected fields
- maintaining a high rate of exchange of qualified human resources
- a dynamic role in the surrounding **innovation** system
- high levels of international visibility and scientific connectivity
- reasonable stability of funding and operating conditions over time



CEITEC's First ERC Grant



Molecular surface rendering of EV71 (human enterovirus 71) virus – Pavel Plevka



European Research Council

Established by the European Commission



CEITEC's ERA Chair Grant - Recruitment

2,7 M Euro for 5 years.

Purpose: To recruit a **transformational figure** in the area of **Chemical and Cellular Biology**



The future ERA Chair will represent the area of research focusing on exploration of cellular systems by advanced imaging methods and their manipulation by means of small organic molecules and/or state-of-the-art physical methods.



CEITEC-EU Life strategic partnership

EU-LIFE mission is to foster excellence, share knowledge, and

influence research policy in life sciences.

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Events of the

NETHERLANDS

INTON WAR UNVERSION

institut**Curie** Descentitie recencies to concer devotes:



7,039 scientists and support personnel 75 running ERC Grants

UNDAÇÃO ALOUSTE GULBENKIAN

IEO

Istituto Europeo di Oncologia

European Institute of Oncology





Miescher Institute **Biomedical Research**



Institute

EITEC

eulife





RG Plant Cytogenomics (Martin Lysak) – recent publications

Mandáková T, Marhold K, **Lysak MA** (2014) The widespread crucifer species *Cardamine flexuosa* is an allotetraploid with a conserved subgenomic structure. *New Phytologist* 201: 982-992.

Cheng F, **Mandáková T**, Wu J, Xie Q, **Lysak MA**, Wang X (2013) Deciphering the diploid ancestral genome of the mesohexaploid *Brassica rapa*. *Plant Cell* 25: 1541-1554.

Mandáková T, Shimizu Inatsugi R, Zozomová-Lihová J, Shimizu K, Kovařík A, Marhold K, **Lysak MA** (2013) The more the merrier: recent hybridization and polyploidy in *Cardamine*. *Plant Cell* 25: 3280-3295.

Long Q, Rabanal FA, Meng D, Huber CD, Farlow A, Platzer A, Zhang Q, Vilhjalmsson BJ, **Mandáková T**, **Lysak MA**, Korte A, Nizhynska V, Voronin V, Korte P, Sedman L, Seren U, Hellmann I, Nordborg M (2013) Massive genomic variation and strong selection in *Arabidopsis thaliana* lines from Sweden. *Nature Genetics* 45: 884-890.

Haudry A et al. (2013) An atlas of over 90,000 conserved noncoding sequences provides insight into crucifer regulatory regions. *Nature Genetics* 45: 891-898. [co-authored by Lysak and Mandáková]

Slotte T et al. (2013) The *Capsella rubella* genome and the genomic causes and the genomic consequences of rapid mating system evolution. *Nature Genetics* 45: 831-835. [co-authored by Lysak and Mandáková]



Comparative chromosome painting: principle



Tribe-Specific Translocation and Karyotype Stasis in the Cardamineae



- Stasis in diploid (2n = 16) and autopolyploid (2n = 32, 48 or 64) genomes
 - Genome reshuffling following allopolyploidy (Leavenworthia, Dentaria)



Seven chromosomes of *Boechera stricta* compared with the eight anestral chromosomes (AK1 - AK8)





Reductions of Chromosome Numbers Were Independent and Used Different Chromosome Breakpoints





Molecular basis of UG-rich RNA recognition by the human splicing factor TDP-43



Structure of the complex reveals novel RNA binding mode

TDP-43 binding to UG-rich RNA associated with **severe forms** of **cystic fibrosis**



Atomic details of the RNA-protein interactions that lead to **aberrant splicing of an essential exon** in severe forms of cystic fibrosis

Lukavsky et al., Nature Structural & Molecular Biology 2013

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Disrupting TDP-43 – UG-rich RNA interactions restores splicing *in vivo*





Small hydrophobic core essential for RNA binding

Alanine mutations abolish RNA binding and **restore splicing** in cystic fibrosis

Searching for small molecule inhibitors:

Setup of fluorescence-based assay for HTS

Lukavsky et al., Nature Structural & Molecular Biology 2013

Thank you for your attention



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